## Supplemental data

Blocking fatty acid-fueled mROS production within macrophages alleviates acute gouty inflammation

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Supplemental Figure 1. MSU crystals activate zebrafish macrophages. (A) Crystal lengths following 18/22 gauge needle dissociation and sonication. (B) Hindbrain expression of *mmp9/krt4* (transverse and dorsal views) within MSU crystal-injected larva (circle, microinjection site). (C) MSU crystal-injected larvae demonstrating 'low' il1b/irg1 expression (insets, magnified views of hindbrain). Hindbrain macrophage and neutrophil within MSU crystal-injected (D) Tg(mpeg1:nfsB-mCherry;lyz:EGFP) larvae (arrow, intracellular crystals). (E) Temporal quantification of crystal phagocytosis, as in D (n=2 larvae). (F) FluoSpheres within hindbrain macrophages (injected into Tg(mpeg1:nfsB*mCherry*) larvae). (G and I) Expression of *il1b* (G) and *irg1* (I) within PBS-, uric acid-, FluoSphere (F.S.)- and calcium pyorophosphate (CPP) crystal-injected larvae. PBS images in G and I are the same as in Figures 1B and 1D, respectively. (H and J) Temporal quantification of *il1b* (H) and *irg1* (J) expression, as detected in G and I, respectively. (K and L) Flow cytometry quantification of neutrophils within Tg(lyz:EGFP) larvae following indicated treatments (K) and MO/CRISPR-Cas9 injections (L), n=~35 larvae/treatment, 5 biological replicates. Untreated sample in panel L is the same as in panel K. Arrows mark illb/irg1 expression in hindbrain. Data for D-J pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. All error bars, means ± SD. \*\*\*\*P<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test. Scale bars, 50  $\mu$ m (B), 100  $\mu$ m (C) and 10  $\mu$ m (D, F).



Supplemental Figure 2. MSU crystal-driven neutrophil recruitment is suppressed in macrophage-depleted larvae. (A) Hindbrain and tail of DMSO- and metronidazole (Mtz.)-treated Tg(mpeg1:nfsB-mCherry) larvae, 18 hours post treatment (hpt). (B and C) Flow cytometry guantification of macrophages (B) and neutrophils (C) from Tg(mpeg1:nfsB-mCherry) (B) and Tg(lyz:EGFP) (C) larvae as treated in A. (D) Flow cytometry quantification of macrophages from Mtz.treated Tg(mpeg1:EGFP) larvae. (E) Hindbrain and tail of liposomomal PBS (L-PBS)- and liposomal clodronate (L-Clod.)-injected Tg(mpeg1:EGFP) larvae, 18 hpi. (F and G) Flow cytometry quantification of macrophages (F) and neutrophils (G) from Tg(mpeg1:EGFP) and Tg(lyz:EGFP) larvae, respectively, as treated in Expression of *il1b* within MSU crystal-injected DMSO- and E. (H) metronidazole(Mtz.)-treated Tg(mpeg1:nfsB-mCherry) larvae and following L-PBS and L-clodronate (L-Clod.) injection. (I) Quantification of *il1b* expression, as detected in H. (J and K) Temporal quantification of neutrophils within the hindbrain of MSU crystal-injected DMSO-Mtz.-treated and Tg(lyz:EGFP;mpeg1:nfsB-mCherry) larvae (J) and in L-PBS- and L-clodronateinjected Tg(lyz:EGFP) larvae (K), n=13-15 larvae/treatment. For flow cytometry, n=5 groups ~30 larvae/treatment. Arrows mark *il1b* expression in hindbrain. Data for H-K pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. All error bars, means ± SD. \*P<0.05; \*\*P<0.01; \*\*\*P<0.001; \*\*\*\*P<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test (B, C, J) and Student's t test (F, G, K). Scale bars, 50  $\mu$ m (A) and 100  $\mu$ m (H).



Supplemental Figure 3. MSU crystal-driven neutrophil recruitment is dependent on macrophage *il1b* expression and Tnfa production. (A) RT-PCR for *il1b* and tnfa from MSU-injected controlMO-, II1bSBMO- and TnfaSBMO-injected larvae (arrowheads, alternatively spliced transcripts). (B) Immunofluorescence detection of neutrophils within the hindbrain of control MO- and II1b SBMO-injected Tg(lyz:EGFP) larvae following MSU crystal injection and MSU crystal-injected DMSO- and Z-VAD-FMK-treated *Tg(lyz:EGFP*) larvae. (C and D) Quantification of neutrophils, as detected in B, for II1b SBMO-injected (C) and Z-VAD-FMKtreated (D) larvae (n=13-15 larvae/treatment). DMSO-MSU samples in panel D are the same as in Figure, 2B/2E/2F. (E) Immunofluorescence of Tnfa within controlMO-, TnfaSBMO- and TnfaATGMO-injected Tg(mpeg1:EGFP) larvae following MSU crystal injection. (F) Quantification of Tnfa, as detected in E (n=15 larvae/treatment). (G) Immunofluorescence detection of neutrophils within the hindbrain of control MO/PBS-injected and control MO-, Tnfa SBMO- and Tnfa ATGMO-injected Tg(lyz:EGFP) larvae following MSU crystal injection. Control MO-MSU image is the same as in panel B. (H and I) Temporal quantification of neutrophils, as detected in G, for Tnfa SBMO-(H) and TnfaATGMO-(I)injected larvae (n=13-15 larvae/treatment). Control MO-MSU samples are the same as in panel C. Data for B-I pooled from 2 independent experiments. All error bars, means ± SD. \*\*P<0.01; \*\*\*P<0.001; \*\*\*\*P<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test. Scale bars, 50 µm (B, E).



Supplemental Figure 4. MSU crystal-driven macrophage activation and neutrophil recruitment is dependent on MyD88. (A) RT-PCR for myd88 from MSU crystal-injected controlMO- and MyD88SBMO-injected larvae (arrowhead, alternatively spliced transcript). (B) Immunofluorescence of MyD88 within controlMO- and MyD88SBMO-injected Tq(mpeq1:EGFP) larvae following MSU crystal injection. (C) Quantification of MyD88, as detected in B (n=15 larvae/treatment). (D) Expression of *il1b* within controlMO-, and MyD88SBMOinjected larvae following MSU crystal injection. (E) Quantification of il1b expression, as detected in D. (F) Immunofluorescence of Tnfa within controlMOand MyD88SBMO-injected Tg(mpeg1:EGFP) larvae following MSU crystal injection. Control MO-MSU image is the same as in Supplemental Figure 3E. (G) Quantification of Tnfa, as detected in F (n=15 larvae/treatment). Control MO-MSU sample is the same as in Supplemental Figure 3F. (H) Immunofluorescence detection of neutrophils within controlMO- and MyD88SBMO-injected *Tg(lyz:EGFP)* larvae following MSU crystal injection. Control MO-MSU image is the same as in Supplemental Figure 3B/3G. (I) Quantification of neutrophils, as detected in H (n=13-15 larvae/treatment). Control MO-MSU samples are the same as in Supplemental Figure 3C/3H/3I. Arrow marks il1b expression in hindbrain. Data for B-I pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. All error bars, means ± SD. \*\*P<0.01; \*\*\*P<0.001; \*\*\*\*P<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test. Scale bars, 50  $\mu$ m (B, H) and 100  $\mu$ m in (D).



Supplemental Figure 5. Blocking NF-κB signaling suppresses MSU crystal-driven macrophage activation and neutrophil recruitment. (A) Expression of *il1b* within MSU-injected DMSO-, and triptolide-treated larvae. DMSO-MSU image is the same as in Figure 3A. (B) Quantification of *il1b* expression, as detected in A. DMSO-MSU sample is the same as in Figure 3B. (C) Immunofluorescence of Tnfa within MSU crystal-injected DMSO- and triptolide-treated Tg(mpeg1:EGFP) larvae. DMSO-MSU image is the same as in Figure 3C. (D) Quantification of Tnfa, as detected in C (n=15 larvae/treatment). DMSO-MSU sample is the same as presented in Figures 1G (3 hpi)/3D. (E) Immunofluorescence detection of neutrophils within MSU crystal-injected DMSOand triptolide-treated Tg(lyz:EGFP) larvae. DMSO-MSU image is the same as in Figure 4A and Supplemental Figure 3B. (F) Quantification of neutrophils, as detected in E (n=13-15 larvae/treatment). DMSO-MSU samples are the same as in Figures, 2B/2E/2F/4B/4C and Supplemental Figure 3D. (G) Immunofluorescence of Tnfa within the hindbrain of MSU crystal-injected DMSO-treated and mpeg1:dnikbaainjected Tg(mpeg1:EGFP) larvae. (H) Quantification of Tnfa, as detected in G (n=15 larvae/treatment). DMSO-MSU sample is the same as in panel D and Figures 1G (3 hpi)/3D. (I) Immunofluorescence detection of neutrophils within the hindbrain of DMSO-treated and mpeg1:dnikbaa-injected Tg(lyz:EGFP) larvae following MSU crystal injection. DMSO-MSU image is the same as in Figure 2D. (J) Temporal quantification of neutrophils, as detected in I (n=13-15 larvae/treatment). DMSO-MSU samples are the same as in panel F, Figure 2B/2E/2F and Supplemental Figure 3D. Arrow marks il1b expression in

hindbrain. All data pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. All error bars, means  $\pm$  SD. \*\**P*<0.01; \*\*\**P*<0.001; \*\*\*\**P*<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test (D, F, H) and Student's *t* test (J). Scale bars, 100 µm in (A) and 50 µm (C, E).



Supplemental Figure 6. Irg1 contributes to MSU crystal-driven macrophage activation and neutrophil recruitment. (A) RT-PCR for irg1 from control MO-, Irg1 SBMO1- and Irg1 SBMO2-injected larvae following MSU crystal injection (arrowheads, alternatively spliced transcripts). (B) Expression of *il1b* within control MO- and Irg1 SBMO2-injected larvae following MSU crystal injection. Control MO-MSU image is the same as in Figure 5A. (C) Quantification of *il1b* expression, as detected in B. Control MO-MSU sample is the same as in Figure 5B. (D) Immunofluorescence of Tnfa within the hindbrain of control MO- and Irg1 SBMO2-injected Tg(mpeg1:EGFP) larvae following MSU crystal injection. Control MO-MSU image is the same as in Figure 5D and Supplemental Figures 3E/4F. (E) Quantification of Tnfa, as detected in D (n=15 larvae/treatment). Control MO-MSU sample is the same as in Figure 5E and Supplemental Figures 3F/4G. (F) Immunofluorescence detection of neutrophils within the hindbrain of control MO- and Irg1 SBMO2-injected Tg(lyz:EGFP) larvae following MSU crystal injection. Control MO-MSU image is the same as in Figure 6A and Supplemental Figures 3B/3G/4H. (G) Temporal quantification of neutrophils, as detected in F (n=13-15 larvae/treatment). Control MO-MSU samples are the same as in Figure 6B and Supplemental Figures 3C/3H/3I/4I. (H) Target sites for Irg1 gRNAs#1/2 and examples of deletions detected by sequencing amplicons generated from individual gRNA/cas9-injected larvae using highlighted primers, compared to gRNA-only-injected larvae (-/+, deletions and additions [in grey], respectively). I, intron; E, exon; red sequence, gRNA targets (PAM sequence underlined). (I) Expression of cxc/8-/1 and cxc/8-/2 within control MO-, Irg1 SBMO1 and Irg1

SBMO2-injected larvae following PBS and MSU crystal injection (as detected by qPCR at 3 hpi, n=25-30 larvae per sample, 3 biological replicates). (J) Macrophage mROS production (white arrow) within the hindbrain of control MOand Irg1 SBMO2-injected *Tg(mpeg1:EGFP)* larvae following MSU crystal injection (MitoSOX signal displayed as a heatmap, warmer colors represents higher levels of mROS). Control MO-MSU image is the same as in Figure 6D. (K) Quantification of macrophage-specific mROS production, as detected in J (n=10 larvae/treatment). Control MO-MSU sample is the same as in Figure 6E. Black arrow marks *il1b* expression in hindbrain. Data for B-G and J,K pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. Error bars, means  $\pm$  SD. \*\*\**P*<0.001; \*\*\*\**P*<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test. Scale bars, 100 µm (B), 50 µm (D, F) and 10 µm (J).



Supplemental Figure 7. JAK/STAT signaling contributes to macrophage activation and neutrophil recruitment in response to MSU crystals. (A) Expression of *irg1* and *il1b* within MSU crystal-injected DMSO-, Stat3 inhibitor peptide(Stat3) IP)- and AG490-treated larvae. (B and C) Quantification of irg1 (B) and il1b (C) expression, as detected in A. (D) Immunofluorescence of Tnfa within the hindbrain of MSU crystal-injected DMSO-, Stat3 IP- and AG490-treated Tg(mpeg1:EGFP) larvae. DMSO-MSU image is the same as in Figure 3C and Supplemental Figure 5C. (E) Quantification of Tnfa, as detected in D (n=15 larvae/treatment). DMSO-MSU sample is the same as in Figures 1G (3) hpi)/3D/5F and Supplemental Figures 5D/5H. (F) Immunofluorescence detection of neutrophils within the hindbrain of PBS-injected and MSU crystal-injected DMSO-, Stat3 IP- and AG490-treated Tg(lyz:EGFP) larvae. PBS image is the same as in Figure 4A. (G) Temporal quantification of neutrophils, as detected in F (n=13-15 larvae/treatment). DMSO-MSU samples are the same as in Figures, 2B/2E/2F/4B/4C/6C and Supplemental Figures 3D/5F/5J. (H) Macrophage mROS production (white arrow) within the hindbrain of MSU crystal-injected DMSO-, Stat3 IP- and AG490-treated Tg(mpeg1:EGFP) larvae (MitoSOX signal displayed as a heatmap, warmer colors represents higher levels of mROS). (I) Quantification of macrophage-specific mROS production, as detected in H (n=10 larvae/treatment). DMSO-MSU sample is the same as in Figure 6F. Black arrows mark *irg1/il1b* expression in hindbrain. All data pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. All error bars, means ± SD. \*P<0.05; \*\*P<0.01; \*\*\*P<0.001;

\*\*\*\**P*<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test. Scale bars, 100  $\mu$ m (A), 50  $\mu$ m (D, F) and 10  $\mu$ m (H).



Supplemental Figure 8. Hindbrain injection of C16:0 or C18:0 elevates MSU crystal-driven macrophage-specific mROS production through a FAO-dependent mechanism. (A) Live imaging of macrophage mROS production within the hindbrain of *Tg(mpeg1:EGFP)* larvae co-injected with MSU crystals and C16:0 or C18:0 with and without etomoxir (Eto.) treatment. (B and C) Quantification of macrophage-specific mROS production, as detected in A, for C16:0 (B) and C18:0 (C) treatments (n=10 larvae/treatment). DMSO-MSU samples are the same as in Figures 6F/8E and Supplemental Figure 7I. All data pooled from 2 independent experiments. All error bars, means  $\pm$  SD. \**P*<0.05; \*\*\*\**P*<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test. Scale bar, 10 µm (A).



Supplemental Figure 9. Exogenous  $H_2O_2$  can rescue MSU crystal-driven macrophage activation following endogenous mROS depletion. (A) Schematic showing delivery of exogenously supplied  $H_2O_2$  to the hindbrain region and ratiometric HyPer imaging of  $H_2O_2$  levels (488/405 nm ratio is displayed as a heat map, with warmer colors representing higher levels of  $H_2O_2$ ) within the hindbrain region of *HyPer* mRNA-injected larvae 5 min before (-5 min post injection, mpi) and after (5 and 90 mpi) injection of 50  $\mu$ M H<sub>2</sub>O<sub>2</sub>. (B) Expression of *il1b* within PBS-injected (with and without co-injected 50 µM H<sub>2</sub>O<sub>2</sub>) and MSU crystal-injected AG490(1 mM)- and etomoxir(Eto., 250 µM)-treated (with and without co-injected 50  $\mu$ M H<sub>2</sub>O<sub>2</sub>) larvae. PBS image is the same as in Figures 1B/7A and Supplemental Figure 1G. (C) Quantification of *il1b* expression, as detected in B. DMSO-MSU sample is the same as in Figure 10E. (D) Immunofluorescence of The within the hindbrain of PBS/H<sub>2</sub>O<sub>2</sub>(50  $\mu$ M)-injected Tg(mpeg1:EGFP) larvae and MSU crystal-injected AG490(1.0 mM)- and etomoxir(250 µM)-treated larvae, with and without co-injected 50  $\mu$ M H<sub>2</sub>O<sub>2</sub>. (E) Quantification of Tnfa, as detected in D (n=15 larvae/treatment). DMSO-MSU sample is the same as in Figures 1G (3 hpi)/3D/5F/7D/11B and Supplemental Figures 5D/5H/7E. Arrows mark il1b expression in hindbrain. Data for B-E pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. All error bars, means ± SD. \*P<0.05; \*\*P<0.01; \*\*\*\*P<0.0001; n.s., not significant, oneway ANOVA, Dunnett's post hoc test. Scale bars, 100  $\mu$ m (A, B) and 50  $\mu$ m (D).







Supplemental Figure 10. Exogenous  $H_2O_2$  can rescue MSU crystal-driven neutrophil recruitment following endogenous mROS depletion. (A) Immunofluorescence detection of neutrophils within the hindbrain of PBS-injected Tg(lyz:EGFP) larvae (with and without co-injected 50  $\mu$ M H<sub>2</sub>O<sub>2</sub>) and MSU crystalinjected AG490(1.0 mM)- and etomoxir(250 µM)-treated larvae (with and without co-injected 50  $\mu$ M H<sub>2</sub>O<sub>2</sub>). PBS image is the same as in Figure 8A. (B and C) Quantification of neutrophils, as detected in A, for AG490/H<sub>2</sub>O<sub>2</sub>- (B) and etomoxir/H<sub>2</sub>O<sub>2</sub>- (C) treatments (n=13-15 larvae/treatment). DMSO-MSU samples Figures, 2B/2E/2F/4B/4C/6C/8B/8C/11D/11E and the same as in are Supplemental Figures 3D/5F/5J/7G. (D) Quantification of neutrophils, as detected in A, for PBS/H<sub>2</sub>O<sub>2</sub>-treatments (n=13-15 larvae/treatment). All data pooled from 2 independent experiments. All error bars, means  $\pm$  SD. \*P<0.05; \*\*P<0.01; \*\*\*P<0.001; \*\*\*\*P<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test. Scale bar, 50  $\mu$ m (A).



Supplemental Figure 11. Phenotypic drug screen uncovers chrysin. piperlongumine and camptothecin as inhibitors of *irg1* expression in activated macrophages. (A) Schematic of drug screen to uncover repurposed drugs that suppress *irg1* expression in activated macrophages. (B) Expression of *irg1* within MSU crystal-injected DMSO-, chrysin-, piperlongumine- and camptothecintreated larvae. DMSO-MSU image is the same as in Supplemental Figure 7A. (C) Quantification of *irg1* expression, as detected in B. DMSO-MSU sample is the same as presented in Supplemental Figure 7B. (D) Time-lapse confocal imaging of macrophage activation (EGFP expression within mCherry<sup>+</sup> macrophages) within the hindbrain of MSU crystal-injected DMSO-, chrysin-, piperlongumineand camptothecin-treated Tg(irg1:EGFP;mpeg1:nfsB-mCherry) larvae. (E-G) Quantification of macrophage activation (measured as fluorescence intensity of EGFP within individual mCherry<sup>+</sup> macrophages), as detected in D, within PBS-, MSU crystal + DMSO-, MSU crystal + chrysin- (E), MSU crystal + piperlongumine-MSU crystal camptothecin-treated (F) and +(G) *Tg(irg1:EGFP;mpeg1:nfsB-mCherry)* larvae. MSU + DMSO and PBS samples are the same in panels E-G. Arrow marks *irg1* expression in hindbrain. Data for B and C pooled from 2 independent experiments. Numbers in parentheses, frequency of larvae with displayed phenotype. Scale bars, 100  $\mu$ m (B) and 50  $\mu$ m (D).



Supplemental Figure 12. Model of Irg1-dependent, FAO/mROS-driven activation of macrophages in response to MSU crystals. Pharmacologic and genetic interventions used in this study to support this model are shown in red.



Supplemental Figure 13. Comparisons between control groups used in this study. (A) Control groups used when assessing the effects of pharmacologic treatments (DMSO), genetic depletion of macrophages (mpeg1:nfsB/DMSO), liposomal clodronate-mediated depletion of macrophages (L-PBS) and MOmediated depletion studies (Control MO) on the temporal recruitment of neutrophils following MSU crystal injection (n=13-15 larvae/treatment). DMSO-MSU samples the in Figures, are same as 2B/2E/2F/4B/4C/6C/8B/8C/11D/11E/13B-D and Supplemental Figures 3D/5F/5J/7G/10B/10C. mpeq1:nfsB/DMSO and L-PBS samples are the same as in Supplemental Figure 2J and 2K, respectively. Control MO samples are the same as in Figure 6B and Supplemental Figures 3C/3H/3I/4I/6G. (B) Control groups used when assessing the effects of pharmacologic treatments (DMSO) and MO-mediated depletion studies (Control MO) on macrophage-specific Tnfa production following MSU crystal injection (n=15 larvae/treatment). DMSO-MSU sample is the same as in Figures 1G (3 hpi)/3D/5F/7D/11B/12D and Supplemental Figures 5D/5H/7E/9E. Control MO-MSU sample is the same as in Figure 5E and Supplemental Figures 3F/4G/6E. (C) Control groups used when assessing the effects of pharmacologic treatments (DMSO) and MO-mediated depletion studies (Control MO) on macrophage-specific mROS production (MitoSOX signal) following MSU crystal injection (n=10 larvae/treatment). DMSO-MSU sample is the same as in Figures 6F/8E/13F and Supplemental Figures 71/8B/8C. Control MO-MSU sample is the same as in Figure 6E and Supplemental Figure 6K. All data pooled from 2 independent experiments. All error bars, means  $\pm$  SD. \*\*\*\**P*<0.0001; n.s., not significant, one-way ANOVA, Dunnett's post hoc test.