

SUPPLEMENTAL DATA

Supplemental Materials and Methods

Cell Culture

THP-1 cells (ATCC; Manassas, VA) were cultured in DMEM (Invitrogen-GIBCO, Grand Island, NY), supplemented with 10% FCS (Hyclone, Logan, UT) and penicillin/streptomycin (P/S; Invitrogen-GIBCO), whereas bone marrow macrophages (BMMs) were cultured in RPMI 1640 (Invitrogen-GIBCO) and 20% L929 cell conditioned media, both supplemented with 10% FCS and P/S, as previously reported [1-3].

Additional Treatments and Assays

Some cells were stimulated with nigericin (20 μ M; EMD-Millipore), Imject Alum Adjuvant (Alum; 200 μ g/ml; Thermo Scientific) and purified *Bacillus anthracis* lethal toxin (LT; lethal factor + protective antigen; a generous gift of J. Mogridge; [4]). Some cells were pretreated with the isopeptidase inhibitor G5 (3,5-bis[(4-Nitrophenyl)methylene]-1,1-dioxide,tetrahydro-4H-thiopyran-4-one; EMD-Millipore, Billerica, MA) that exhibits specificity for BRCC3 [5], caspase 1 inhibitor YVAD (Enzo Life Sciences, Farmingdale, NY) and PKR inhibitor (2-Aminopurine; Sigma-Aldrich; [6]), as indicated.

HMGB1 in culture supernatants was measured by immunoblotting with anti-HMGB1 (Cell Signaling; Beverly, MA) and IL-18 was measured by ELISA (R&D Systems).

Bacterial Infections

Overnight cultures of *S. typhimurium* SL1344 [7] were diluted 1:50 and grown to an $OD_{600} = 0.6$ in LB. After washing in PBS, bacteria were resuspended in Optimem (10^9 CFU/ml; Invitrogen-GIBCO) and added (MOI = 25) to LPS primed BMMs. One hour later cultures were washed with PBS and grown in Optimem supplemented with gentamycin (100 μ g/ml; 5h). *L. pneumophila* (JR32; restriction-defective Philadelphia-1) and flagellin minus (Fla-) strains (JR32 background)

were grown in AYE broth or on CYE plates [1]. Previously characterized strains ectopically over expressing *cdgS3*, *cdgS3B* or *cdgS4* were placed on a flagellin null background [8]. BMMs were infected with post-exponential phase, IPTG (isopropyl- β -D-thiogalactopyranoside; 1 mM, 6h; Sigma Aldrich) treated *L. pneumophila* (MOI = 10), as previously reported [1].

TMRM & mitoSox staining

Mitochondrial reactive oxygen species (mROS) were measured by FACS (LSR Fortessa; BD Biosciences) in mitoSox (Invitrogen; 2.5 μ M, 20 min in Optimem) stained BMMs. FACS or fluorescence microscopy (Zeiss Axiovert 200M; Oberkochen, Germany) was used to measure mitochondrial membrane tetramethyl rhodamine methyl ester (TMRM; Anaspec, Fremont, CA; 200 nM, 30 min in Optimem) uptake (i.e., $\Delta\psi_m$). FACS data were analyzed with FlowJo software (Tree Star; Ashland, OR).

Table S1. Primer Sequences

Mu-IFN- β	Forward 5' TCC AAG AAA GGA CGA ACA TTC G 3'
	Reverse 5' TGA GGA CAT CTC CCA CGT CAA 3'
Hu-ASC	Forward 5' CTG CTG GAT GCT CTG TAC 3'
	Reverse 5' GCT GGT GTG AAA CTG AAG 3'
Hu-NLRP3	Forward 5' GTT CAC TGC CTG GTA TCT 3'
	Reverse 5' GTG CTT GCC ATC TTC ATC 3'
Hu-MPYS	Forward 5' ACT TAC AAT CAG CAT TAC AAC A 3'
	Reverse 5' GTC AGC CAT ACT CAG GTT 3'
Mu-MPYS	Forward 5' GCC TGG TCA TAC TAC ATT G 3'
	Reverse 5' GCA GCA TAT CTC GGA ATC 3'
Mu-GAPDH	Forward 5' CAT CAA GAA GGT GGT G 3'
	Reverse 5' CCT GTT GCT GTA GCC 3'
β -actin	Forward 5' GCT CCT CCT GAG CGC AAG T 3'
	Reverse 5' TCG TCA TAC TCC TGC TTG CTG AT 3'

Figure S1. Additional inflammatory responses in murine BMMs

(A) Secreted IL-1 β was measured by ELISA in LPS primed BMMs (left panel) treated with ATP (5 mM; 6h) or transfected (X-tremegene HP; Xtrm) with c-diGMP (10 nmol; 6h), c-diAMP (10 nmol; 6h), cGMP (10 nmol; 6h) or GTP (10 nmol; 6h), as in Figure 1A. PMA differentiated THP-1 cells (right panel) were treated with ATP (5 mM; 6h) or transfected (Lipofectamine 2000; lpftm) with c-diAMP (cdA; 5nmol; 6h), c-diGMP (cdG; 5 nmol; 6h) or poly(dA:dT) (dA:dT; 5 μ g/ml; 6h), as Figure 1A. Graphs present means \pm standard error from three independent experiments. (B) Secreted IL-1 β was measured by ELISA in LPS primed BMMs treated with KCl, Glibenclamide (Gli), 2-APB or U73122 prior to stimulation with Imject Alum Adjuvant (Alum; 200 μ g/ml; Thermo Scientific) or nigericin (20 μ M; 1h), as in Figure 2A. Graphs present means \pm standard error from two independent experiments.

Figure S2. c-diGMP stimulation of cells deficient of inflammasome components.

(A) Secreted IL-1 β was measured by ELISA in LPS primed C57Bl/6J wild type (WT), NLRP3 $[-/-]$ (NLRP3), or ASC $[-/-]$ (ASC) BMMs treated as in Figure 3. Graphs present means \pm standard error from three independent experiments. (B) IL-1 β was measured by ELISA in LPS primed C57Bl/6J (Bl6) WT and AIM2 $[-/-]$ treated as in panel A. Graphs present means \pm standard error from two independent experiments. (C) IL-1 β was measured by ELISA in LPS primed 129 WT and NLRP1b $[-/-]$ BMMs treated with ATP (5 mM; 6h), *B. anthracis* (LT; 1 μ g/ml, as indicated), or transfected with c-diGMP (cdG;10 nmol; 6h) and c-diAMP (cdA; 10 nmol; 6h), as in panel A. Graphs present means \pm standard error from two independent experiments. (D) IL-1 β levels were measured by ELISA in LPS primed BMMs, which were pretreated with the PKR inhibitor, 2 amino purine (2-AP; 1 mM, 30 min) prior to stimulation with ATP (5 mM; 30 min) or transfection (X-tremegene HP; Xtrm) with c-diGMP (10 nmol; 6h). Graphs present means \pm standard error from three independent experiments. (E) IL-18 levels were measured by ELISA on supernatants from panel C. Graphs present means \pm standard error from three independent experiments. (F)

Expression of ASC (left panel) and NLRP3 (right panel) was evaluated in different THP-1 cell lines by Q-PCR, as indicated. Results were normalized to β -Actin and reported as relative fold change. Graphs present means \pm standard error from three independent experiments.

Figure S3. c-diGMP stimulation is not associated with mitochondrial perturbation.

Mitochondrial ROS production (**A**) and mitochondrial membrane potential (**C**) were evaluated by FACS in LPS primed BMMs stimulated with ATP (5 mM, 6h), c-diGMP (cdG; 10 nmol; 6h) or poly(dA:dT) (dA:dT; 5 μ M; 6h), after staining with mitoSox (2.5 μ M; 20 min) and TMRM (200 nM; 30min), respectively. (**B, D**) Mean fluorescent intensity (MFI) quantification of the data from panels A & C, respectively. Data are representative of 3 independent experiments. (**E**) Mitochondrial membrane potential after TMRM staining was also evaluated by fluorescence microscopy on BMMs cells treated as in panel C.

Figure S4. HMGB1 production and Deubiquitinylation in c-diGMP stimulated BMMs.

(**A**) Secreted HMGB1 was measured by immunoblotting (anti-HMGB1; Cell Signaling) supernatants collected from C57Bl/6J BMMs treated with ATP, c-diGMP and c-diAMP, as in Figure 1A. (**B**) Secreted IL-1 β was measured by ELISA in LPS primed C57Bl/6J BMMs treated with YVAD (50 μ M; 30 min) or G5 (3,5-bis[(4-Nitrophenyl)methylene]-1,1-dioxide,tetrahydro-4H-thiopyran-4-one; 1 & 2 μ M; 30 min) prior to treatment with ATP, c-diGMP and c-diAMP, as in Figure S1. Graphs present means \pm standard error from three independent experiments.

Figure S5. IL-1 β is required for c-diGMP mediated neutrophil recruitment in vivo.

(**A**) Peritoneal fluid recovered from mice described in Figure 4 was stained with antibodies to Ly6G and F4/80 and evaluated by FACS. Data are from individual mice, representative of two independent experiments (n=6 mice for c-di-GMP and n=4 mice for PBS treatment). (**B**) Peritoneal fluid was evaluated by FACS analysis 16 h after intraperitoneal injection of PBS or 200

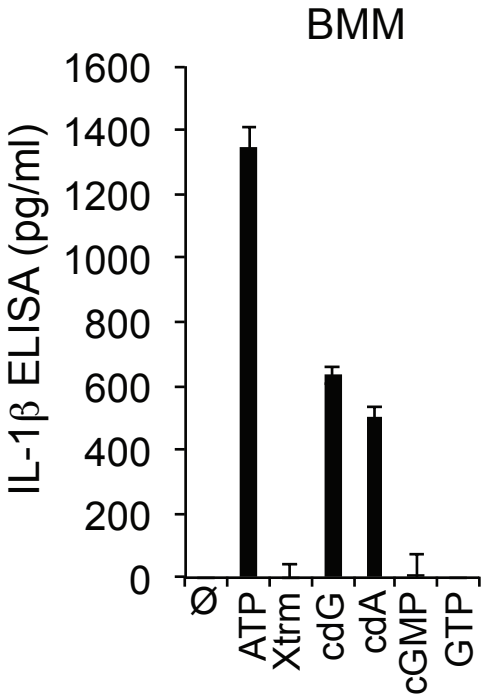
nmol of c-diGMP (cdG) as detailed in Figure 4. In another study 40 µg of anti-IL-1β antibody were injected (IP; 200 µl PBS) 30 min prior to c-diGMP injection (cdG + α-IL-1β). The histogram reports the number of neutrophils recruited to the peritoneum.

References

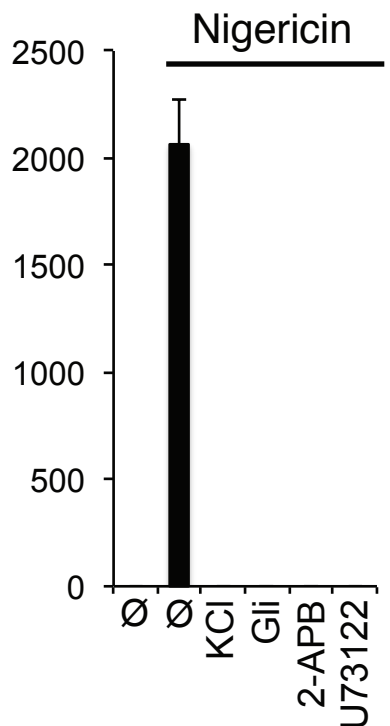
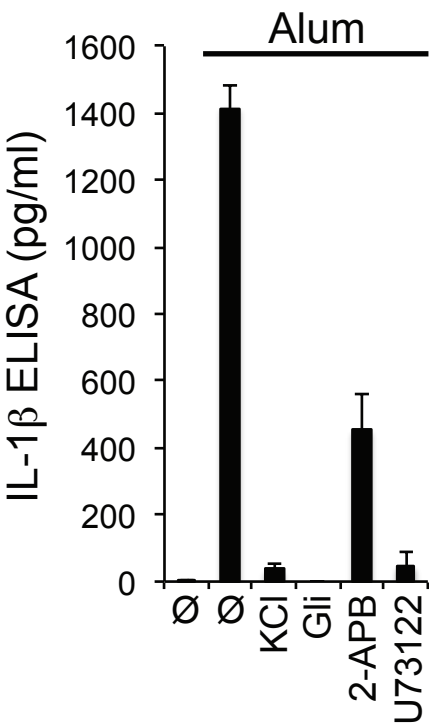
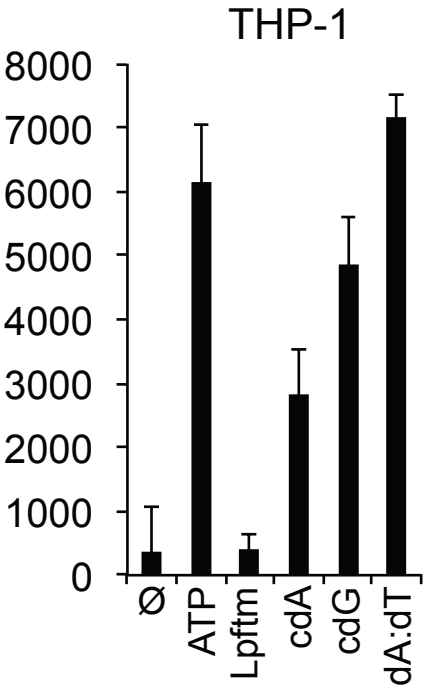
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Figure S1, Abdul-Sater et al.

A



B



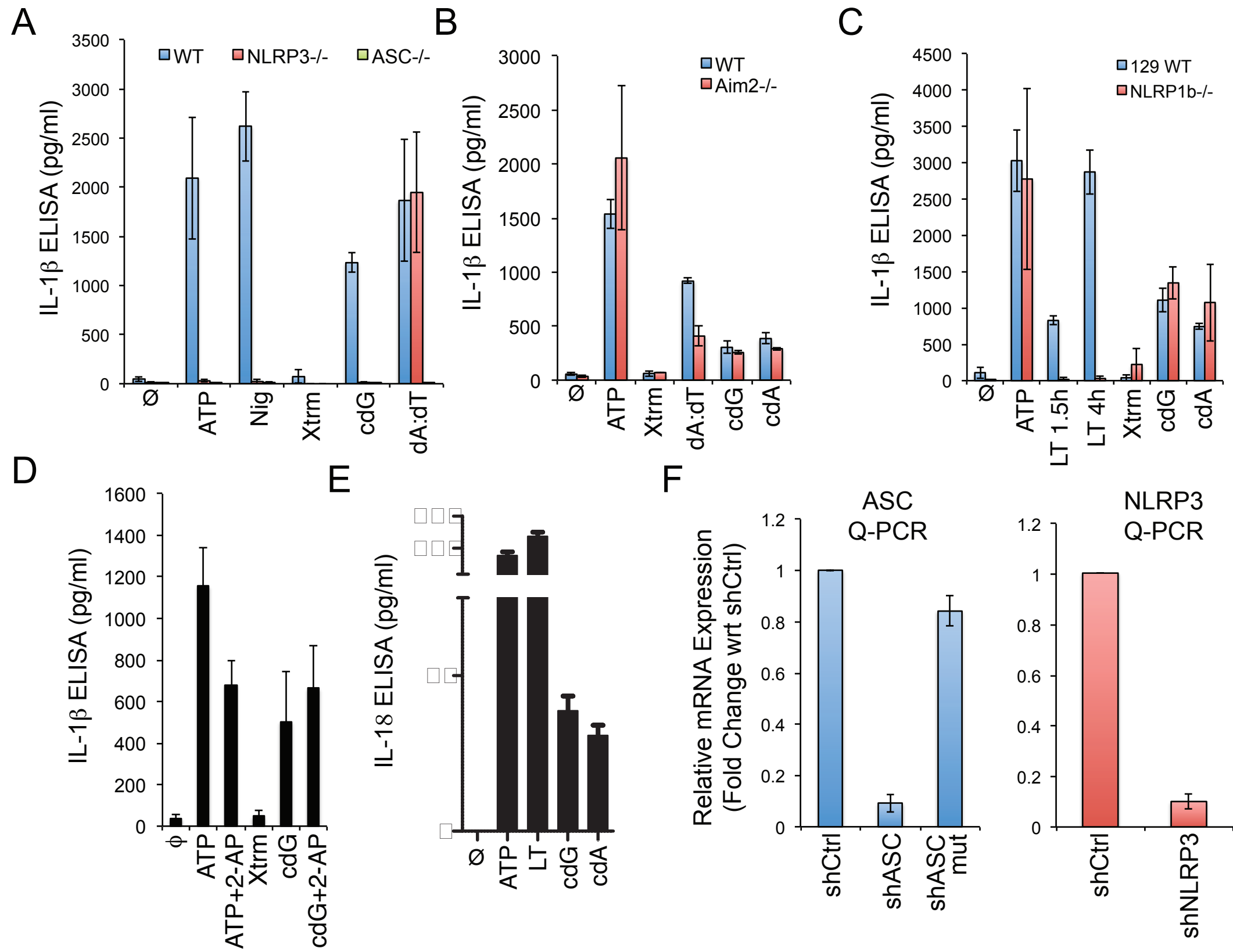
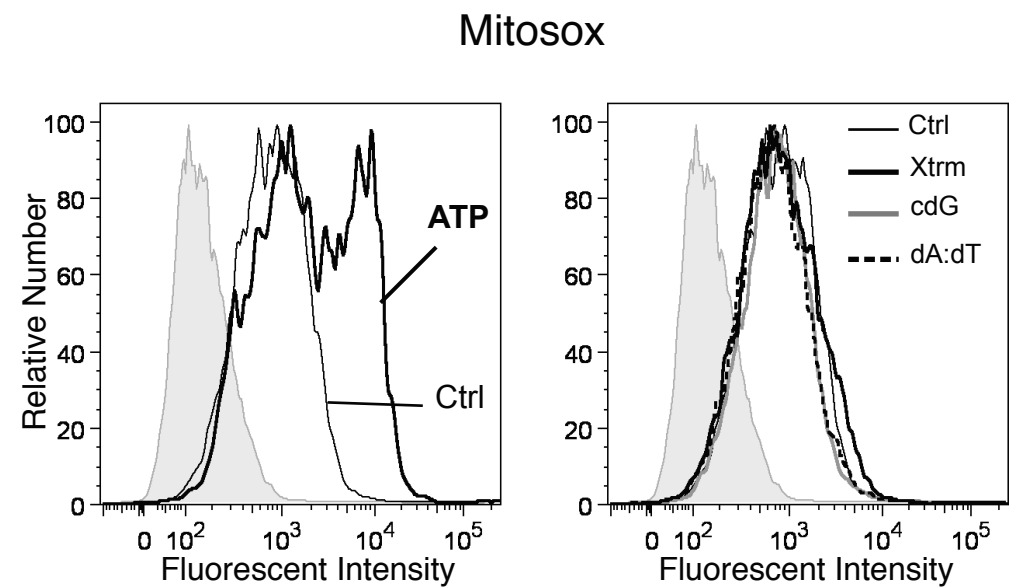
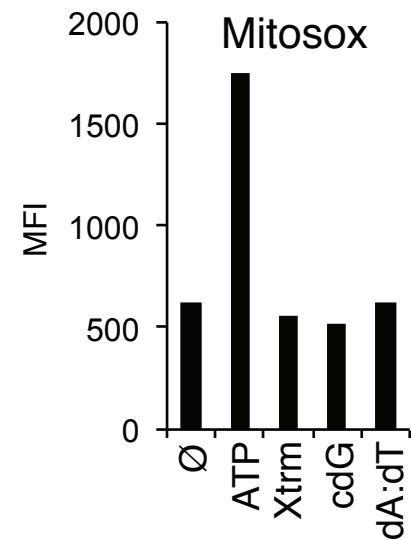


Figure S3, Abdul-Sater et al.

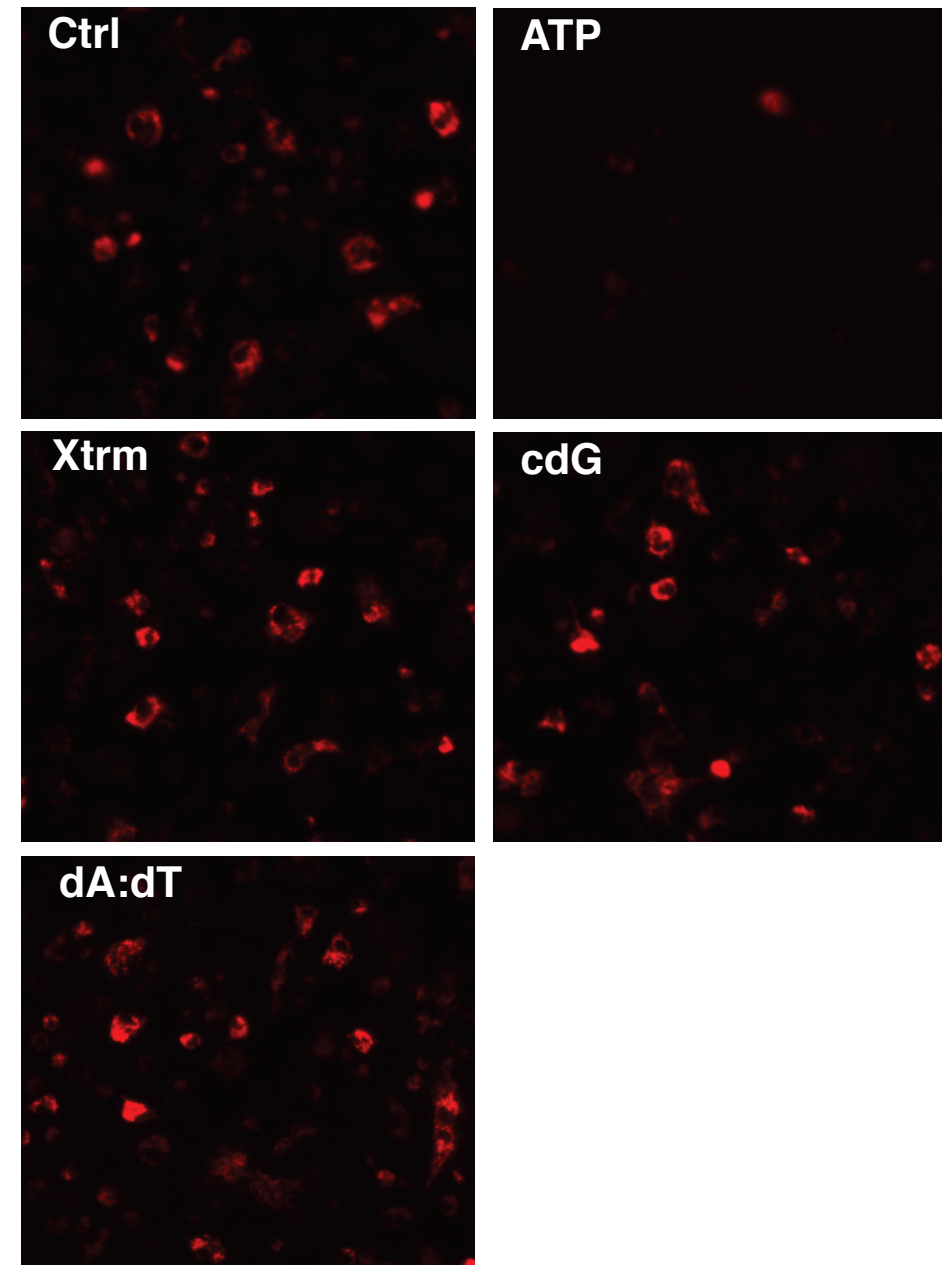
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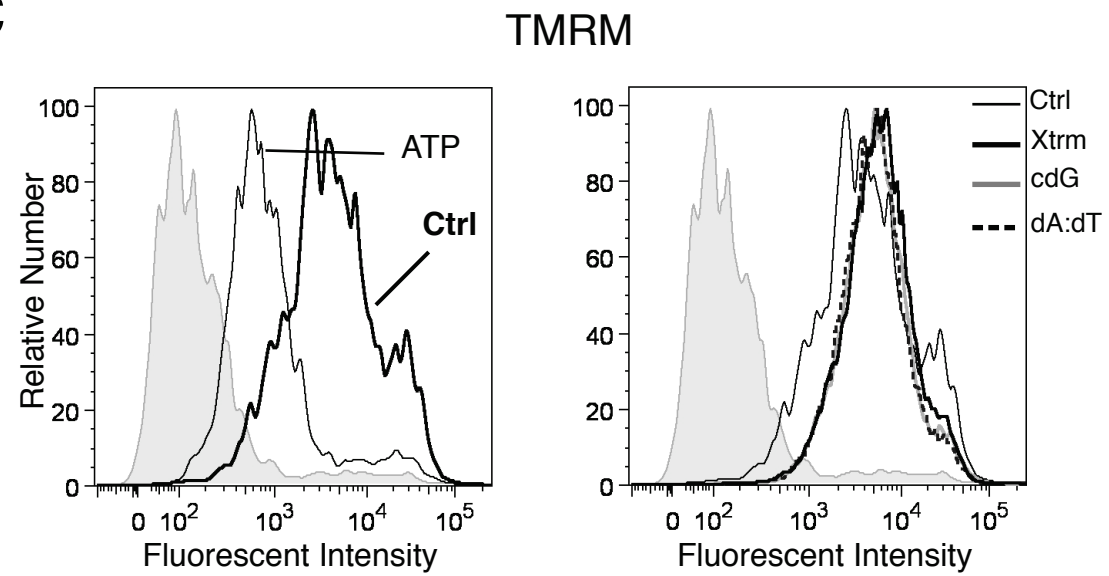
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E



C



D

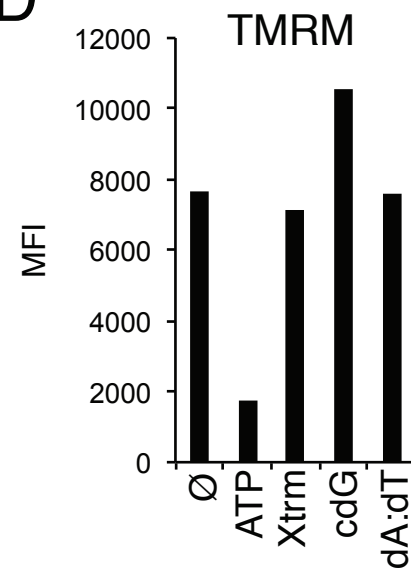


Figure S4, Abdul-Sater et al.

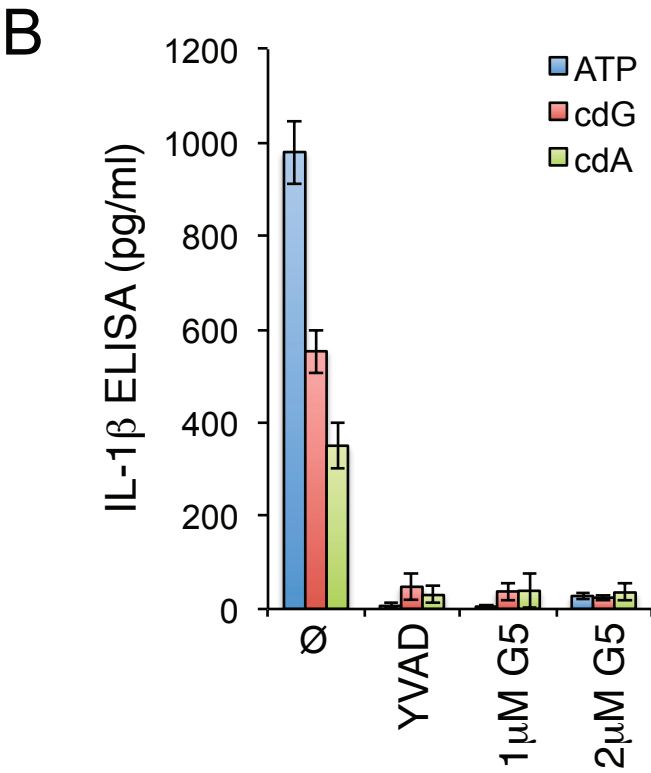
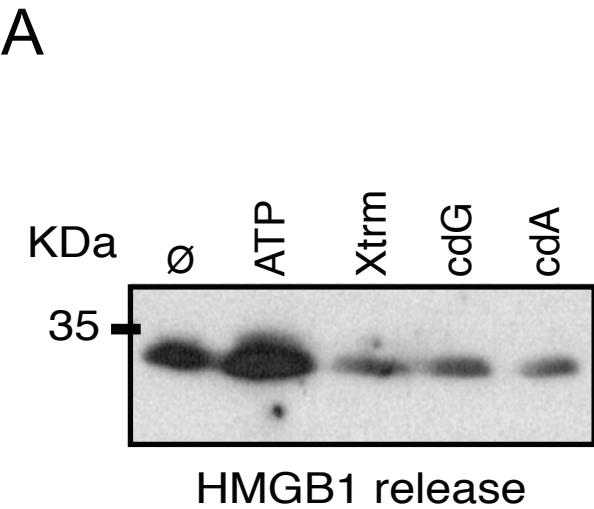
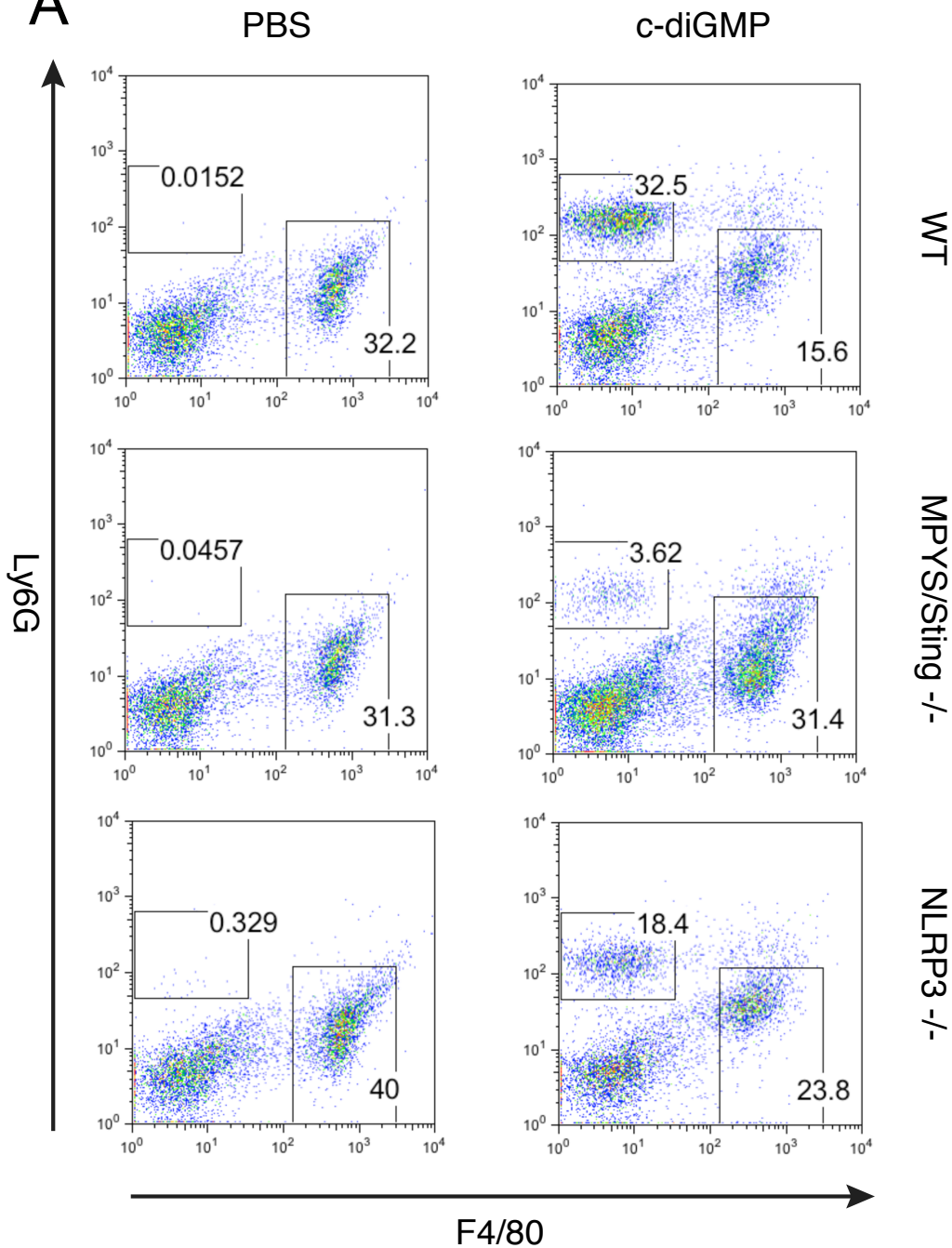


Figure S5, Abdul-Sater et al.

A



B

