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# *Trichomonas vaginalis* Induces NLRP3 Inflammasome Activation and Pyroptotic Cell Death in Human Macrophages

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## Abstract

*Trichomonas vaginalis* is a sexually transmitted, eukaryotic parasite that causes trichomoniasis, the most common nonviral, sexually transmitted disease in the USA and worldwide. Little is known about the molecular mechanisms involved in the host immune response to this widespread parasite. Here we report that *T. vaginalis* induces NLRP3 inflammasome activation in human macrophages, leading to caspase-1 activation and the processing of pro-IL-1β to the mature and bioactive form of the cytokine. Using inhibitor-based approaches, we show that NLRP3 activation by *T. vaginalis* involves host cell detection of extracellular ATP via P2X$_7$ receptors and potassium efflux. In addition, our data reveal that *T. vaginalis* inflammasome activation induces macrophage inflammatory cell death by pyroptosis, known to occur via caspase-1 cleavage of the gasdermin D protein, which assembles to form pores in the host cell membrane. We found that *T. vaginalis*-induced cytolysis of macrophages is attenuated in gasdermin D knockout cells. Lastly, in a murine challenge model, we detected IL-1β production in vaginal fluids in response to *T. vaginalis* infection in vivo. Together, our findings mechanistically dissect how *T. vaginalis* contributes to the production of the proinflammatory IL-1β cytokine and uncover pyroptosis as a mechanism by which the parasite can trigger host macrophage cell death.

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## Introduction

*Trichomonas vaginalis* is a eukaryotic parasite that infects the urogenital tract of women and men. The parasite is transmitted through sexual intercourse and, after having colonized its human host, causes the disease trichomoniasis, which is the most common nonviral, sexually transmitted infection (STI) in the USA [1] and worldwide [2]. While *T. vaginalis* STI can be asymptomatic in many individuals [3], others experience inflammation of the vagina, cervix, or prostate [4]. The inflammatory response mounted against the parasite is predicted to fuel multiple

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adverse health effects associated with *T. vaginalis* infection. These include a higher incidence of premature births [5], an increased risk of cervical [6, 7] or aggressive prostate cancer [8, 9], and an increased risk of acquiring [10] and potentially transmitting the human immunodeficiency virus (HIV) to a sexual partner [11].

Despite the serious public health threat posed by *T. vaginalis*, limited knowledge exists regarding the molecular mechanisms by which *T. vaginalis* elicits inflammation. As an extracellular parasite, *T. vaginalis* survives in the human body via uptake of nutrients from host cells to which it can adhere and/or phagocytose and lyse, including vaginal or prostate epithelial cells [12], and red or white blood cells [13]. Currently, human clinical data regarding the specific innate immune response to *T. vaginalis* infections are scarce, and existing knowledge largely stems from human coinfection studies or from in vitro experiments. Cauci and Culhane [14] reported a significant increase in vaginal IL-1β levels in pregnant women who had bacterial vaginosis and a *T. vaginalis* coinfection compared to pregnant women with only bacterial vaginosis. Indeed, inflammatory profiling of human monocytes and monocyte-derived macrophages in response to *T. vaginalis* infection identified IL-1β as an upregulated cytokine [15–17]. However, the cellular mechanisms and requirements for the production of this important immune response factor during macrophage and *T. vaginalis* encounters are unknown. Here we delineate the cellular pathway that is activated in macrophages by *T. vaginalis* leading to the maturation and release of IL-1β.

Inflammasomes are multimeric cytosolic complexes that sense the presence of pathogens or changes in cellular homeostasis and activate commensurate innate immune responses. The inflammasome complex is defined by the responding sensor protein, which has characteristic structural domains and includes the nucleotide-binding oligomerization domain-like receptors (NLRs) NLRP1, NLRP3, and NLRC4, as well as the AIM2 and pyrin proteins [18]. Inflammasome assembly is initiated by NLR recognition of pathogen-associated molecular patterns (PAMPs) or endogenous danger-associated molecular patterns (DAMPs) [18]. Upon detecting these stimuli, the sensor proteins oligomerize and associate with the adaptor protein ASC in a multiprotein complex that recruits pro-caspase-1 and promotes its autoproteolytic processing [19, 20]. Mature protease caspase-1 then cleaves the pro-forms of inflammatory cytokines IL-1β and IL-18 to generate the mature bioactive forms capable of binding their cognate receptors and initiating cell signaling [21–27]. Two major outcomes of inflammasome activation

are processing and release of mature IL-1β and induction of a rapid cell death termed pyroptosis [28, 29].

In this study, we show that *T. vaginalis* activates NLRP3 inflammasomes in human macrophages, leading to bioactive IL-1β production and pyroptotic cell death. The parasite-driven NLRP3 inflammasome activation involves sensing of ATP as a DAMP and potassium efflux, and IL-1β release in response to *T. vaginalis* can be detected in vivo in a mouse vaginal challenge study. Thus, *T. vaginalis* activation of NLRP3 inflammasomes leads not only to a strong macrophage proinflammatory response against the parasite, consistent with recent findings in prostate epithelial cells [30], but also to elimination of the macrophages in the process. The balance of innate immune detection and inflammatory response is likely to influence *T. vaginalis* colonization and pathogenesis differentially depending on the magnitude and stage of infection.

## Materials and Methods

### Growth of Cells and Culture

*T. vaginalis* strains RU393 (New York, NY, USA; ATCC 50142 [31]) and MSA1132 (Mt. Dora, FL, USA [12]) were both obtained from Patricia Johnson's Laboratory at UCLA. Parasites were treated with 50 μg/mL chloramphenicol and 5 μg/mL tetracycline for 6 days and then frozen down as stocks. The parasites were thawed from these stocks and passaged daily for less than 2 weeks, as previously described [32], in the continued presence of chloramphenicol and tetracycline. Human THP-1 monocytes (ATCC TIB-202) were cultured in RPMI medium supplemented with 10% fetal bovine serum (FBS), 0.05 mM 2-mercaptoethanol, 0.2% D-glucose, 10 mM HEPES, and 1 mM sodium pyruvate. The THP-1 monocytes were differentiated to macrophages by treating the cells with 25 nM PMA (phorbol 12-myristate 13-acetate; Thermo Fisher Scientific) for 24 h.

### THP-1 Cas9 Cell Line

A Cas9 stable cell line was generated by transducing THP-1 cells with Lenti-Cas9-Blast lentivirus (Addgene 52962-LV). Spinfection was performed by spinning down THP-1 cells with viral particles supplemented with 10 μg/mL polybrene at 1,000 *g*, 32 °C, for 1 h. After 48 h, cells were selected for stable expression of *Streptococcus pyogenes* Cas9 using blasticidin (10 μg/mL). Ten days after selection, the cells were harvested and Cas9 expression was confirmed by immunoblotting and qPCR.

### Cloning of GSDMD sgRNAs

Two sgRNA sequences previously shown to successfully target the gasdermin D (GSDMD) gene were cloned into the lentiGuide-Puro vector (Addgene; Plasmid #52963) as described previously [33]. The sgRNA sequences were as follows: GSDMD KO1 5′-TGAGTGTGGACCCTAACACC-3′ [from 34] and GSDMD KO2 5′-CTTGCTTTAGACGTGCAGCG-3′ [from 33; GeCKOv2 Library, identifier #HGLibA_20413].

### GSDMD Knockout Cell Lines

Lentivirus was packaged in HEK 293T cells (ATCC CRL-3216) using the Fugene HD transfection reagent (Promega) and 4 μg of the gasdermin D sgRNA-lentiGuide-Puro construct, 4 μg of the packaging plasmid psPAX2 (Addgene; Plasmid #12260), and 4 μg of the VSV-G envelope expressing plasmid pMD2-G (Addgene; Plasmid #12259). THP-1 Cas9 cells (described above) were spinfected with 10 μg/mL polybrene and viral particles for 1 h at 1,000 $g$, 32 °C. After 48 h, cells were selected for stable expression of the sgRNAs using puromycin (1 μg/mL). After 10 days of selection, cells were cloned by limiting dilution. Two clonal cell lines (KO1 and KO2), one from each sgRNA, were harvested and gasdermin D knockout was confirmed by immunoblotting assays and PCR analysis.

### T. vaginalis *Co-Culture Experiments*

A total of $1 \times 10^5$ THP-1 cells were seeded and differentiated in 96-well plates. *T. vaginalis* RU393 parasites were spun down at 2,061 $g$, washed once, and then resuspended in RPMI + 2% heat-inactivated FBS. Serial dilutions were performed to attain the desired *T. vaginalis*-to-macrophage ratios. Medium was aspirated from the THP-1 macrophages and 100 μL of *T. vaginalis* cell suspension added to each well. For inhibitor or chemical treatment experiments, parasites were premixed with the vehicle or test compound and then added to the macrophages.

### Cytokine and Caspase-1 Protein Analysis

IL-1β cytokine levels and caspase-1 protein levels present in cell culture supernatants were quantified using the DuoSet IL-1β ELISA kit and the Quantikine Human Caspase-1/ICE ELISA Kit (both from R&D Systems).

### IL-1β Signaling Assay

HEK-Blue™ IL-1β Cells (InvivoGen) were purchased and grown per the company's specifications. A total of 50,000 IL-1β Sensor Cells were seeded per well of a 96-well plate; 50 μL of cell supernatant from each experimental sample or a recombinant human IL-1β (InvivoGen) dilution was added to the IL-1β Sensor Cells and incubated at 37 °C, 5% $CO_2$, for 18 h. Secreted alkaline phosphatase activity was assessed by adding 50 μL of the HEK-Blue IL-1β reporter cell supernatants onto 150 μL of QUANTI-Blue™ reagent (InvivoGen) and measuring the optical density at 620 nm using an EnSpire Plate Reader (PerkinElmer). Background levels from THP-1 only or THP-1 cells treated with vehicle were subtracted from all values to report IL-1β signaling levels above background levels.

### Caspase-1 Activity Assay

Caspase-1 activity was assessed using the Caspase-1/ICE Colorimetric Assay Kit (BioVision). A total of $2 \times 10^6$ THP-1 macrophages were seeded and differentiated in 6-well plates. *T. vaginalis* RU393 parasites were spun down at 2,061 $g$, washed once, and resuspended in RPMI + 2% heat-inactivated FBS; 1.5 mL of parasite suspension were added to each well. After a 2.5-h co-incubation, the cell supernatants were removed and the cells lysed with cell lysis buffer. Caspase-1 activity was assessed per the manufacturer's specifications in whole cell lysates normalized by equal protein amounts.

### Reagents and Inhibitors

*Escherichia coli* LPS, ATP, KCl, NaCl, and adenosine 5′-triphosphate, periodate oxidized sodium salt (oxATP) were purchased from Sigma. The Ac-YVAD-CMK caspase-1 peptide inhibitor was purchased from Enzo, and the NLRP3/AIM2 inhibitor CRID3 was purchased from Tocris.

### Filter Experiment Assays

Co-incubation experiments were performed in the presence of a Millicell®-96 Cell Culture Insert Plate PCF with a pore size of 0.4 μm (Millipore). For contact conditions, *T. vaginalis* cell suspensions were placed under the filter to allow direct interaction between *T. vaginalis* and THP-1 cells. For filter conditions that prevented physical contact, *T. vaginalis* was placed on top of the filter.

### Lactate Dehydrogenase Cytolysis Assays

Cytolysis was assessed by assaying for lactate dehydrogenase (LDH) release from cells using the CytoTox-ONE™ Homogeneous Membrane Integrity Assay Kit (Promega). The manufacturer's specifications were used to perform the assays and calculate the percent cytolysis compared to 100% lysis controls. For cytolysis analysis of gasdermin D knockout cell lines, cell lysis was compared to THP-1 Cas9 parental cells (THP-1 wild type).

### Immunoblot Analysis

A total of $2 \times 10^6$ THP-1 macrophages were seeded and differentiated per well of a 6-well plate. After co-incubation with *T. vaginalis*, the cells were washed 2 times with PBS and lysed with RIPA Buffer (Thermo Fisher Scientific) containing HALT™ protease inhibitor cocktail. Samples were separated by SDS-PAGE, immunoblotted, and visualized with SuperSignal® WestPico Chemiluminescent Substrate (Thermo Fisher Scientific) and CL-XPosure™ Film (Thermo Fisher Scientific). The following antibodies were used for immunoblotting: anti-IL-1β (AF-201-NA; Cell Signaling), anti-β-actin (A5316; Sigma), anti-gasdermin D (20770-1-AP; Proteintech), anti-goat IgG HRP-conjugated (HAF017; R&D), and ECL Amersham anti-mouse IgG HRP-linked (NA931; GE Healthcare).

### Animal Experiments

Six-week-old female C57BL/6 mice were purchased from The Jackson Laboratory. The mice were pretreated by daily intraperitoneal (i.p.) injections with 100 μL of 2 mg/mL Dexamethasone Sodium Phosphate Injection, USP (West-Ward Pharmaceuticals) for 4 days prior to infection. The day before infection, the mice were also injected i.p. with 100 μL of 5 mg/mL β-estradiol resuspended in sesame oil (both from Sigma). For animal infections, *T. vaginalis* MSA1132 strain was washed twice and resuspended in sterile DPBS (Dulbecco's phosphate-buffered saline; Corning). A total of $6 \times 10^6$ *T. vaginalis* in a final volume of 10–15 μL were used to infect the mice vaginally. Three days after infection, vaginal lavages were collected by washing the vaginal lumen 4 times with 50 μL of DPBS (200 μL total) using Gel-Loading pipet tips (Thermo Fisher Scientific).

### Statistical Analysis

For the in vitro experiments, statistical significance was calculated using a Student $t$ test. For IL-1β detection in the animal studies, Grubbs' test was performed to test for the presence of experi-
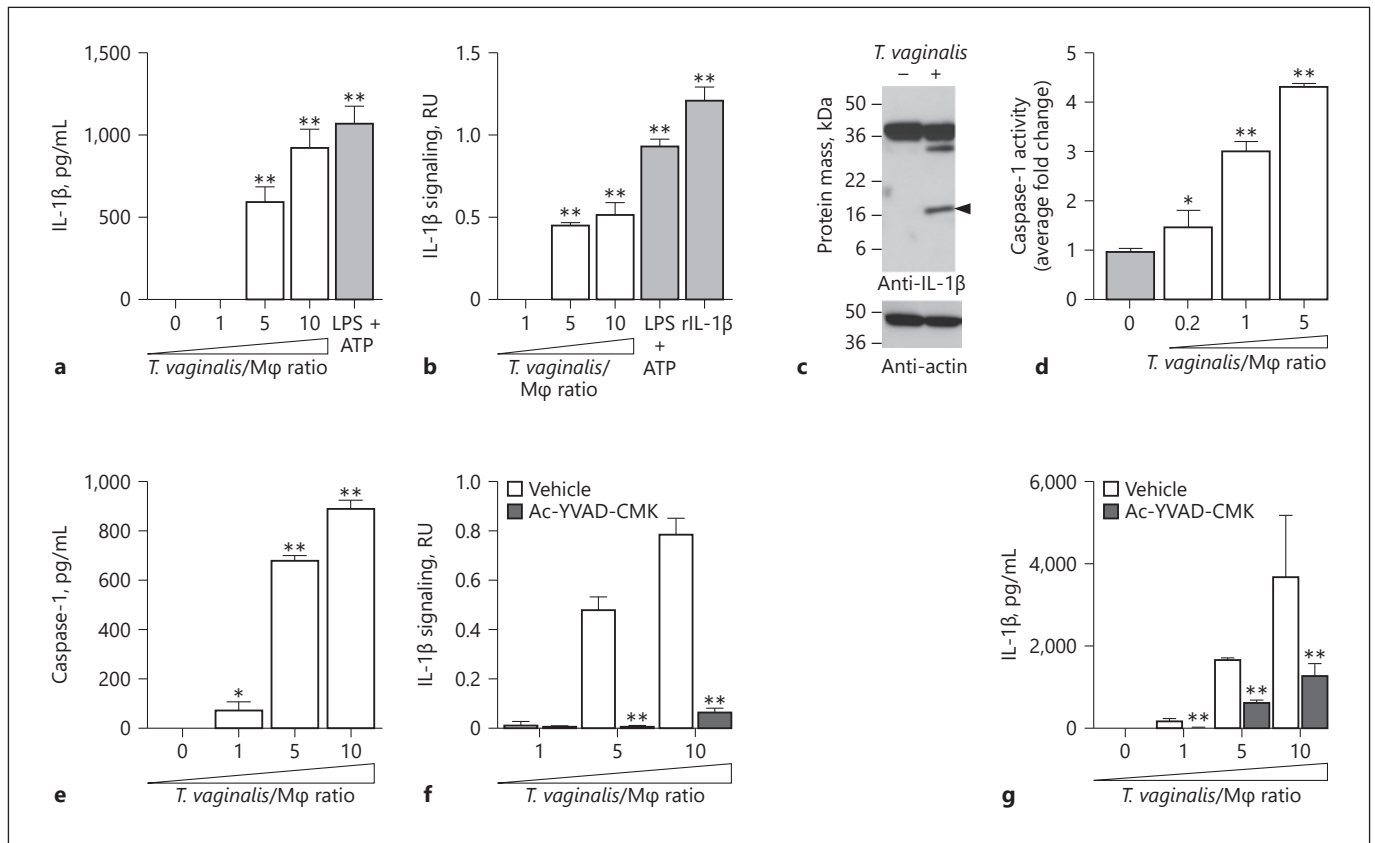
**Fig. 1.** THP-1 exposure to *Trichomonas vaginalis* leads to bioactive IL-1β production via caspase-1 activity. THP-1 macrophages were incubated with *T. vaginalis* using different multiplicities of infection (MOIs; *T. vaginalis*:host cell) for 4 h unless otherwise stated. As a positive control for inflammasome activation, THP-1 cells were stimulated with 10 ng/mL LPS + 5 mM ATP. To test for the presence of processed IL-1β, experiment supernatants were placed on HEK-Blue IL-1β reporter cells. Bioactive IL-1β binds to the IL-1 receptor on IL-1β reporter cells, activating production of a reporter enzyme. Reporter enzyme activity was quantified spectrophotometrically and is proportional to IL-1β signaling activity, shown as relative units (RU) after subtracting background readings from supernatants of THP-1 cells alone or THP-1 cells treated with vehicle control. As a positive control, recombinant bioactive IL-1β (rIL-1β) was added to the HEK-Blue IL-1β reporter cells. **a** IL-1β in cell supernatants was measured by ELISA. **b** Bioactive IL-1β was detected using the HEK-Blue IL-1β reporter cells. **c** Immunoblot detection of the IL-1β cleavage product (17 kDa; arrowhead) in whole cell lysates of THP-1 macrophages exposed to *T. vaginalis* for 4 h at an MOI of 1. Full-length pro-IL-1β (31 kDa) could be detected in both samples. Actin loading control and molecular weight markers in kilodalton are also shown. **d** THP-1 macrophages were exposed to *T. vaginalis* for 2.5 h. The amount of caspase-1 activity was assayed by measuring cleavage of the p-nitroanilide-labeled YVAD peptide (YVAD-pNA) spectrophotometrically. The fold change relative to THP-1 cells alone is shown. **e** Amounts of caspase-1 protein released into cell supernatants as quantified by ELISA. **f**, **g** THP-1 cells were co-incubated with *T. vaginalis* in the presence of the caspase-1 peptide inhibitor Ac-YCAD-CMK or vehicle control. Levels of bioactive IL-1β were detected using the IL-1β reporter cells (**f**) and IL-1β protein levels were detected by ELISA (**g**). For each experiment, samples were assayed at least in triplicate. Representative results from 2 (**d**, **e**) or 3 (**a–c**, **f**, **g**) independent experiments are shown. The graphs show the mean with standard deviations. * $p < 0.05$, ** $p < 0.01$, compared to THP-1 cells (**a**, **b**, **d**, **e**) or vehicle treatment at that MOI (**f**, **g**).

mental outliers. One cytokine value from the uninfected mouse group (777.768 pg/mL) and 1 value from the *T. vaginalis*-infected group (266.183 pg/mL) were identified as outliers. A Mann-Whitney test was used to test for statistical significance across the remaining samples. $p < 0.05$ was considered as statistically significant.

## Results

### T. vaginalis *Stimulates Bioactive IL-1β Secretion in a Caspase-1-Dependent Manner*

The NLRP3 inflammasome is highly expressed and best characterized in cells of the myeloid cell lineage in-
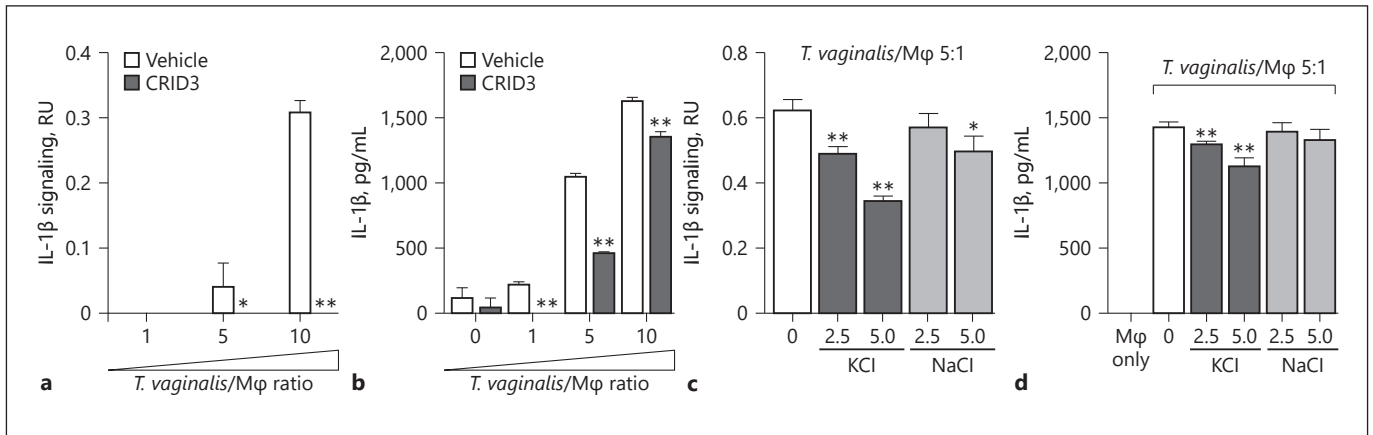
**Fig. 2.** *Trichomonas vaginalis* activates NLRP3 inflammasomes in THP-1 macrophages. THP-1 macrophages were exposed to *T. vaginalis* at a multiplicity of infection of 5 (*T. vaginalis*:host) for 4 h in the presence of CRID3 (**a**, **b**), an NLRP3/AIM2 inflammasome inhibitor, or exogenous KCl and NaCl (**c**, **d**). Levels of bioactive IL-1β were detected using the IL-1β reporter cells (**a**, **c**) and total IL-1β protein levels were detected by ELISA (**b**, **d**). A representative result from 3 independent experiments is shown. The graphs show the mean with standard deviations. * $p < 0.05$, ** $p < 0.01$, compared to THP-1 cells co-incubated with *T. vaginalis* vehicle treatment.

cluding macrophages and dendritic cells [35, 36]. As macrophages are among the most abundant immune effector cells in the female reproductive tract [37], we investigated whether *T. vaginalis* activates NLRP3 inflammasomes in human macrophages. For our studies, we used the human THP-1 monocyte-like cell line, which is differentiated to macrophages using PMA treatment. NLRP3 inflammasome activation involves two signals. Signal 1, known as the priming step, occurs after receptor engagement, and signal transduction leads to NF-κB activation, which in turn stimulates increased transcription of the genes encoding pro-IL-1β and NLRP3 [38–40]. Signal 2, the activation step, is initiated by diverse microbial PAMPs and host cell DAMPs which lead to the physical assembly of the NLPR3 inflammasome complex. Examples of NLRP3 signal 2 activators include bacterial surface proteins, microbial pore-forming toxins, and viral RNA, as well as host cell DAMPs such as extracellular ATP, amyloid-β, and monosodium urate crystals [41, 42].

We found a dose-dependent increase in IL-1β secretion from THP-1 macrophages at increasing multiplicities of infection (MOIs) of *T. vaginalis* (Fig. 1a); LPS and ATP, two well-described NLRP3 inflammasome inducers that provide signal 1 and signal 2, respectively [43], served as positive controls. While IL-1β protein has been detected by ELISA in the monocyte/macrophage cytokine response to *T. vaginalis* exposure [15–17], it was unclear if this represented the pro-form of IL-1β released secondary to parasite-induced cell lysis or whether it was the pro-

cessed and bioactive form of the inflammatory cytokine. To test the bioactivity of IL-1β generated upon THP-1 macrophage exposure to *T. vaginalis*, we added cell supernatants from the co-culture experiments to HEK-Blue IL-1β Sensor Cells (InvivoGen). These reporter cells express IL-1R (IL-1 receptor) to initiate cell signaling and trigger the production of a SEAP reporter enzyme that can convert a colorimetric substrate to provide a readout for IL-1β signaling. With recombinant bioactive IL-1β serving as a positive control for IL-1β signaling activity, Figure 1b shows that *T. vaginalis* co-incubation with THP-1 macrophages led to processing of pro-IL-1β and generation of bioactive IL-1β capable of IL-1R-mediated cell signaling. LPS and ATP served as positive controls for NLRP3 inflammasome activation in the THP-1 cells. The immunoblot analysis in Figure 1c confirms detection of the 17-kDa bioactive IL-1β cleavage product in whole cell lysates from THP-1 macrophages exposed to *T. vaginalis*, which was not detected in the uninfected control. Conversely, the full-length pro-IL-1β protein, which has a molecular weight of 31 kDa but migrates with an apparent molecular weight of ~36 kDa, was detected in both uninfected THP-1 macrophages and macrophages exposed to *T. vaginalis*. Therefore, macrophage co-incubation with *T. vaginalis* leads to active production of processed and bioactive IL-1β.

Canonical inflammasome assembly and activation leads to the autoproteolytic processing of the proenzyme form of caspase-1 (45 kDa) to its active form composed

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Tsai/Nizet

of a heterodimeric tetramer of the p20 and p10 subunits [44]. To quantify caspase-1 activity, we assayed cleavage of the chromogenic peptide YVAD-pNA. After 2.5-h co-incubation with *T. vaginalis*, we detected a ~4-fold increase in caspase-1 activity from THP-1 cells at an MOI of 5 and a ~3-fold increase at an MOI of 1 (Fig. 1d). Caspase-1 can be released into cell supernatants after inflammasome activation [45], and we found a dose-dependent release of high amounts of caspase-1 in THP-1 supernatants following exposure to increasing MOIs of *T. vaginalis* (Fig. 1e). Treatment with the specific Ac-YVAD-CMK peptide inhibitor of caspase-1 markedly ablated bioactive IL-1β production (Fig. 1f), a result corroborated when total IL-1β was quantified by ELISA (Fig. 1g). Together, these results indicate that upon exposure to *T. vaginalis*, caspase-1 activity is increased in THP-1 cells and leads to the processing of pro-IL-1β to bioactive IL-1β.
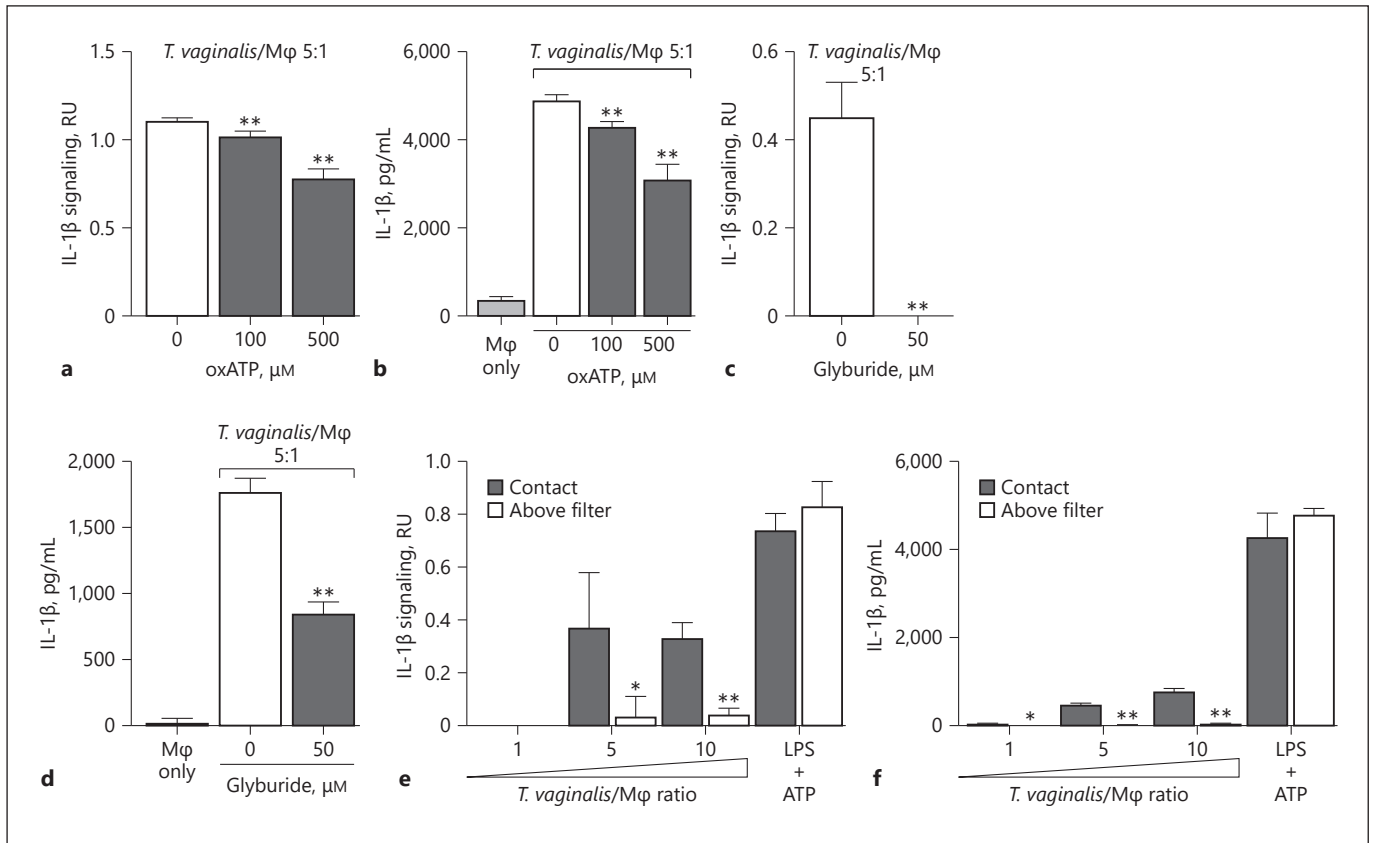
### T. vaginalis *Activates NLRP3 Inflammasomes*

Multiple innate immune sensor proteins can be assembled to form different types of inflammasome complexes [18]. Many inflammasomes are activated by bacterial components such as LPS or flagellin [46] that are not found in the eukaryotic *T. vaginalis* parasite. NLRP3 inflammasomes were strong molecular candidates for the observed generation of bioactive IL-1β, since they can also be activated in response to host cell perturbations and by sensing DAMPs potentially released during cellular injury inflicted by *T. vaginalis*. To investigate the role of NLRP3 as the sensor protein driving IL-1β processing, we co-incubated *T. vaginalis* with THP-1 cells in the presence or absence of CRID3, which inhibits ASC oligomerization to block both NLRP3 and AIM2 inflammasomes [47]. CRID3 treatment completely ablated bioactive IL-1β signaling from THP-1 macrophages exposed to *T. vaginalis* at an MOI of 5 or 10 (Fig. 2a); similar reductions in total IL-1β detected by ELISA were observed at each MOI (Fig. 2b). A variety of ligands or cell perturbations can serve as upstream activators of NLRP3 inflammasome activation; however, the precise molecular mechanism shared by these various pathways to result in NLRP3 inflammasome activation is still a subject of active investigation. $K^+$ efflux has been identified as a common signaling event in many cases of NLRP3 inflammasome activation [41, 48, 49]. Therefore, we compared the levels of IL-1β signaling in THP-1 cells co-incubated with *T. vaginalis* in the presence or absence of added extracellular KCl to counteract the effects of $K^+$ efflux [49] and observed a dose-dependent inhibition of IL-1β signaling (Fig. 2c) and total IL-

1β protein levels (Fig. 2d). As a control for addition of extracellular ions, exogenous addition of NaCl only slightly modified IL-1β signaling (Fig. 2c) and had no effect on IL-1β protein levels (Fig. 2d). Together these results indicate that *T. vaginalis* induction of IL-1β secretion in THP-1 macrophages is mediated through NLRP3 inflammasome activation in a scenario where $K^+$ efflux is required.

### ATP-P2X₇ *Receptor Signaling and Macrophage Contact Contributes to* T. vaginalis-*Induced Inflammasome Activation*

Extracellular ATP serves as a DAMP by binding to cell surface $P2X_7$ receptors that provoke $K^+$ efflux from the cell and subsequent NLRP3 inflammasome activation [50–55]. To test whether ATP plays a role in inflammasome activation by *T. vaginalis*, we co-incubated the parasites with THP-1 macrophages in the presence of ox-ATP, which covalently and irreversibly binds to P2 family receptors (P2XR), inhibiting their ATP-mediated ion transport and ATP-dependent IL-1β release [54, 56]. oxATP treatment led to a dose-dependent reduction in IL-1β signaling and IL-1β protein levels in THP-1 macrophages exposed to *T. vaginalis* (Fig. 3a, b). To further probe $P2X_7$ receptor involvement, we used glyburide (also known as glibenclamide), which inhibits $P2X_7$ receptors and is also a broad-spectrum inhibitor of ATP-binding cassette transporters. As a result of blocking $K^+$ efflux, glyburide inhibits NLRP3 inflammasomes [57]. Glyburide treatment completely ablated the production of bioactive IL-1β in *T. vaginalis*-infected THP-1 macrophages (Fig. 3c), and more than halved the amount of IL-1β protein detected by ELISA (Fig. 3d). Together, these results indicate that exposure of macrophages to *T. vaginalis* leads to inflammasome activation via $P2X_7$ receptor-mediated $K^+$ efflux in response to extracellular ATP.

To determine if direct contact between the parasite and the THP-1 cells was necessary to induce inflammasome activation, we placed a filter in between the macrophages and *T. vaginalis*, which allows soluble materials to pass through but prohibits direct contact between the parasites and the THP-1 macrophages. In conditions where *T. vaginalis* was placed above the filter (no direct contact), the secretion and processing of IL-1β were markedly reduced (Fig. 3e, f). Thus, physical contact of the host macrophages with the parasites is critical for sensing either an unknown PAMP on *T. vaginalis* or, more likely, a DAMP (e.g., ATP) arising from the host cell generated upon cell lysis that serves as a signal for inflammasome activation.

**Fig. 3.** Response to extracellular ATP via P2X$_7$ receptor signaling and physical contact contributes to *Trichomonas vaginalis* inflammasome activation. THP-1 macrophages were exposed to *T. vaginalis* at a multiplicity of infection (MOI) of 5 (*T. vaginalis*:host) for 4 h in the presence of oxidized ATP (oxATP; **a**, **b**) or glyburide (**c**, **d**), an ATP-gated P2X$_7$ receptor inhibitor. **e**, **f** A 0.4-µm membrane filter that allowed soluble materials to flow through but prevented contact was placed in between THP-1 macrophages and *T. vaginalis* in the "above filter" condition, or parasites were placed below the filter to allow contact between *T. vaginalis* and macrophages ("contact" condition). THP-1 cells were treated with 10 ng/mL LPS + 5 mM ATP as a positive control for NLRP3 activation. Levels of bioactive IL-1β were detected using the IL-1β reporter cells (**a**, **c**, **e**) and total IL-1β protein levels were detected by ELISA (**b**, **d**, **f**). Representative results from 3 independent experiments are shown in **a–d**, and from 2 independent experiments in **e** and **f**. The graphs show the mean with standard deviations. **a–d** ** $p < 0.01$ compared to THP-1 cells exposed to *T. vaginalis* in the presence of vehicle control. **e**, **f** * $p < 0.05$, ** $p < 0.01$, compared to the contact with *T. vaginalis* condition at that MOI.

### T. vaginalis *Inflammasome Activation Leads to THP-1 Macrophage Cell Death by Pyroptosis*

In addition to IL-1β secretion, NLRP3 inflammasome activation can lead to an inflammatory cell death called pyroptosis. Cell lysis occurs after the mature caspase-1 protease has cleaved the gasdermin D protein [28], which then oligomerizes [58] and forms pores in the plasma membrane [29], prompting cell swelling and membrane rupture. Pyroptosis is recognized as a distinct form of cell death in large part due to its fast nature (compared to apoptosis) and its accompanying "fiery" inflammatory response [59]. Pyroptosis is commonly detected by monitoring downstream membrane damage that leads to re-

lease of intracellular proteins like LDH. Of note, LDH assays have been the standard assay used by many investigators to monitor *T. vaginalis* lysis of vaginal epithelial cells and prostate epithelial cells [12]. To test how *T. vaginalis*-induced inflammasome activation may influence macrophage pyroptosis, we examined the effect of inflammasome inhibitors on macrophage LDH release following infection with the parasite. Co-incubation of cells in the presence the caspase-1 inhibitor Ac-YVAD-CMK led to a significant reduction in host cell lysis at each MOI (Fig. 4a). Similar reductions in THP-1 macrophage lysis by *T. vaginalis* were also obtained with the NLRP3/AIM2 inflammasome inhibitor CRID3 (Fig. 4b). A complete re-
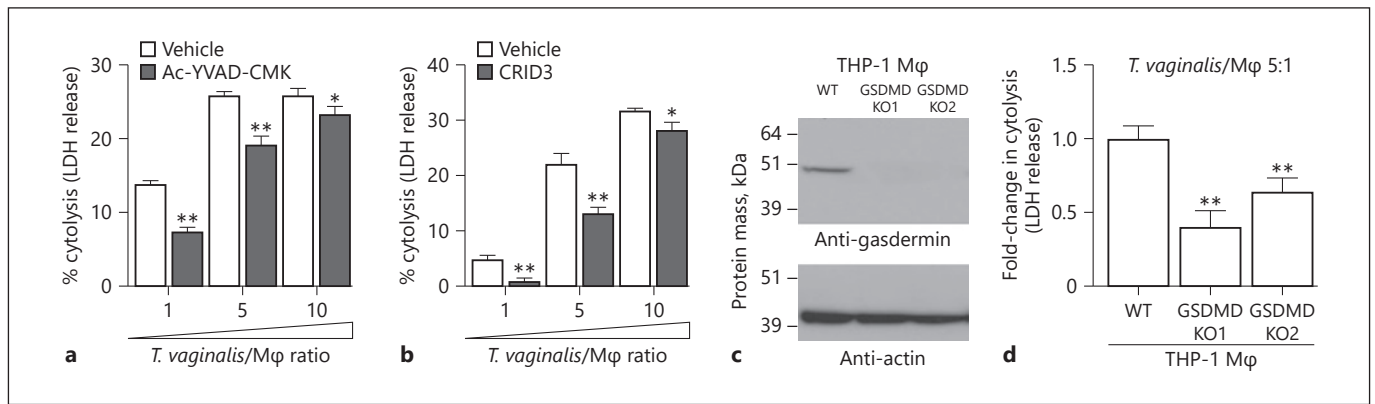
**Fig. 4.** Inflammasome activation by *Trichomonas vaginalis* contributes to pyroptosis. Cytolysis of host cells was assessed by measuring lactate dehydrogenase (LDH) release from dying macrophages. **a, b** THP-1 macrophages were incubated with *T. vaginalis* using different multiplicities of infection (MOIs; *T. vaginalis*:host cell) for 4 h in the presence of vehicle, the caspase-1 peptide inhibitor Ac-YVAD-CMK (**a**), or the NLRP3/AIM2 inhibitor CRID3 (**b**). The data are representative from 3 experiments. The graphs show the mean with standard deviations. * $p < 0.05$, ** $p < 0.01$, compared to vehicle treatment at that MOI. **c, d** Gasdermin D (GSDMD) was knocked out in THP-1 cells using CRISPR-Cas9. **c** Confirmation of gasdermin D knockout (KO) in 2 clones by Western blot analysis using an anti-gasdermin D antibody. Actin loading control and molecular weight markers in kilodalton are also shown. **d** GSDMD KO cells or wild-type (WT) THP-1 cells were exposed to *T. vaginalis* at an MOI of 5 for 4 h. The average fold change in percent cytolysis compared to THP-1 WT cells from 3 independent experiments with standard errors of the mean are shown. ** $p < 0.01$ compared to THP-1 WT cells.

duction in cell death is not unexpected, as inhibition of pyroptosis can activate apoptosis [60], and caspase-1 peptide inhibitors may yield incomplete inhibition of LDH release after inflammasome activation [20, 61]. Therefore, to further investigate the induction of pyroptosis by *T. vaginalis* at a molecular level, we generated two independent knockouts of gasdermin D in THP-1 macrophages using CRISPR-Cas9 as confirmed by Western blot (Fig. 4c) and PCR evidence of genome editing. When the gasdermin D knockout macrophages were exposed to *T. vaginalis*, we observed a 40–60% reduction in LDH release (cell death) compared to wild-type THP-1 cells (Fig. 4d). These results reveal that *T. vaginalis* can kill host macrophages through pyroptosis.

As a pilot study to determine if activation of IL-1β release is observed in vivo, we modified a previously established mouse model of *T. vaginalis* infection [62]. Unfortunately, *T. vaginalis* mouse models have required estrogen treatment together with the synthetic glucocorticoid and immunosuppressant dexamethasone [63] prior to and after *T. vaginalis* infection in order to facilitate colonization by the parasite [62]. To investigate inflammasome activation by *T. vaginalis* under minimal immune suppression conditions, we inoculated mice with one of the most virulent clinical strains available, called MSA1132 [12], using an optimized protocol illustrated schemati-

cally in Figure 5a. *T. vaginalis* strain MSA1132 also yielded bioactive IL-1β production in our in vitro studies (data not shown). Three days after parasite or mock (PBS) infection, we collected vaginal lavages and examined IL-1β protein levels by ELISA. *T. vaginalis* infection led to a statistically significant increase in IL-1β secretion in vivo (Fig. 5b), consistent with our in vitro results.

## Discussion

In this work we show that upon contact with THP-1 human macrophages, *T. vaginalis* can activate NLRP3 inflammasomes, lead to release of bioactive IL-1β, and induce pyroptotic cell death. We thus infer that direct contact of *T. vaginalis* with macrophages is necessary to deliver both signals required for NLRP3 inflammasome activation. Inflammasome activation by *T. vaginalis* is mediated in part by sensing extracellular ATP via $P2X_7$ receptors, which lead to $K^+$ efflux. Midlej and Benchimol [64] have shown that *T. vaginalis* inflicts membrane damage upon host cell contact, a likely requisite for the release of ATP and other DAMPs that can serve as signal 2 for inflammasome activation. 100 μM oxATP can inhibit the ability of 5 mM ATP to stimulate macrophage IL-1β production [65]. Similarly, we observed inhibition of IL-1β
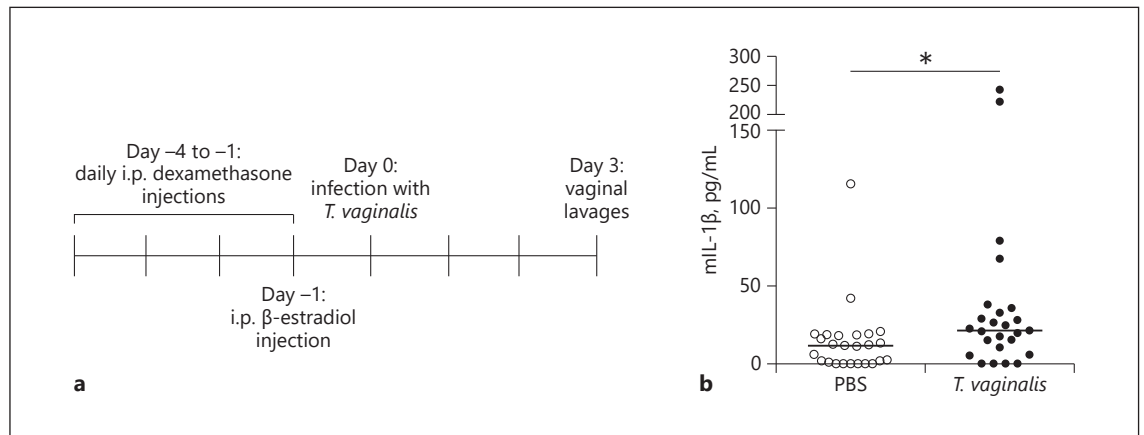
**Fig. 5.** *Trichomonas vaginalis* infection leads to IL-1β production in a mouse model of infection. **a** The timeline shows the pretreatment strategy for female C57BL/6 mice prior to intravaginal *T. vaginalis* inoculation or PBS mock infection. **b** Concentrations of IL-1β protein levels in vaginal lavages collected 3 days after infec-tion quantified by ELISA. The data shown are from 3 independent experiments combined ($n = 25$ mice per treatment group). The median and range are shown. * $p < 0.05$ compared to the PBS uninfected control.

signaling with 100 μM oxATP, but higher doses of oxATP yielded still greater inhibition, so it is likely that significant ATP quantities are generated locally by *T. vaginalis*-induced cell damage allowing macrophage P2X$_7$ receptor activation. We hypothesize that the general mechanism of host cell injury by *T. vaginalis* and subsequent release of DAMPs may underpin our preliminary observation of *T. vaginalis*-induced IL-1β secretion in our murine short-term challenge model.

Several groups have reported *T. vaginalis* activation of NF-κB in vaginal epithelial cells, human monocyte-derived macrophages, murine macrophages, and THP-1 macrophages [16, 17, 66, 67], establishing a precedence for the parasite's ability to provide a signal 1 required for NLRP3 inflammasome priming. To date, the only *T. vaginalis* cell surface component shown to contribute to NF-κB activation is the parasite's glycocalyx [68], now known as Tv lipoglycan [69]. Toll-like receptor (TLR) 2 contributes to sensing of *T. vaginalis* by murine macrophages leading to NF-κB signaling [67]. It remains to be mechanistically investigated what other TLRs contribute to sensing *T. vaginalis*. TLR4 involvement has been implicated by indirect evidence, as cervicovaginal lavages from *T. vaginalis*-infected women stimulated TNF-α production from TLR4-responsive murine splenocytes at higher levels than splenocytes nonresponsive to TLR4 ligands [70]. In HeLa cells, TLR2, TLR4, and TLR9 expression is increased upon infection with *T. vaginalis* [71], and TLR4 upregulation has also been reported in a prostate stromal cell line exposed to the parasite [72]. Further dissection of

how *T. vaginalis* contact with human macrophages leads to the generation and delivery of both signal 1 and signal 2 is a subject of ongoing investigation in our laboratory.

Our work has also demonstrated that *T. vaginalis* activation of the NLRP3 inflammasome in macrophages contributes to their physical lysis via pyroptosis, the first direct demonstration that the parasite can kill host cells through this inflammatory cell death pathway. Gu et al. [30] found that *T. vaginalis* can activate NLRP3 inflammasomes in a prostate epithelial cell line, but measures of cell viability, pyroptosis, or gasdermin D involvement were not reported. Nevertheless, our results and that of Gu et al. [30] add *T. vaginalis* to the growing number of sexually-transmitted pathogens that lead to inflammasome activation [73, 74]. Studying the outcomes of inflammasome activation by STIs is of clinical importance, as the proinflammatory microenvironment in the reproductive tract may affect susceptibility to other STIs or facilitate coinfections [73, 74], and *T. vaginalis* is already associated with gonorrhea, chlamydia, syphilis, and herpes simplex virus types 1 and 2, as well as with an increased risk and transmission of HIV [10, 11, 75]. While our work was performed with THP-1 cells, an established cell line previously used for the study of vaginal colonizing bacteria [76, 77], future experimentation with our murine model and human primary vaginal macrophages is warranted to fully evaluate how *T. vaginalis* activates inflammasomes in the vaginal mucosal environment.

Bioactive IL-1β can exert effects on virtually all cell types and promote a multitude of functional outcomes in

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innate immunity and the shaping of adaptive immunity [78, 79]. These functions include inducing fever, promoting increased leukocyte recruitment to the infection site, and enhancing effector functions and cell survival [78]. Increased neutrophil numbers are reported in vaginal lavages of *T. vaginalis*-infected women [14, 80], but it remains unknown whether in human infections the increased leukocyte recruitment contributes to controlling the infection or to inducing pathology. An example of the latter adverse effects occurs with the extracellular fungus *Candida albicans*, in which NLRP3 inflammasome activation contributes to pathogenesis by promoting increased neutrophil influx and vaginitis in a murine model of infection [81, 82].

The physical lysis of host macrophages through pyroptosis likely mitigates the magnitude of the host innate immune response to *T. vaginalis*. For example, although we detected increased IL-1β production upon *T. vaginalis* infection in our murine infection model, we did not observe any gross visible phenotypes. Similarly, a majority of infected individuals are also asymptomatic [3]. This dichotomy highlights the complexity in the biology of *T. vaginalis* infections, in which inflammasome activation occurs in response to the parasite; however, the lysis of IL-1β-producing cells by the parasite may affect the amounts and effects of the bioactive IL-1β produced. The parasite is also highly motile with 5 flagella [83, 84]. This property likely influences how long the parasite remains in contact with host cells to allow for parasite sensing, how effective released host effector molecules are at targeting the parasite, and the effect of recruited leukocytes, if the parasite can physically swim away. Other parasite virulence factors also help dampen the local immune response – and potentially the effects of inflammasome activation on adaptive immunity. For example, *T. vaginalis* proteases can degrade complement components and antibodies [85], and the parasite can also lyse B cells and T cells [15]. *T. vaginalis* also expresses ectoATPases which can degrade extracellular ATP [86], the latter of which we identified to serve as a DAMP for NLRP3 inflammasome activation. Future work testing the effect of parasite mutants and uncovering such parasite virulence factors will allow us to broaden our understanding of the overall innate immune response, of its impact on *T. vaginalis* survival in the host, and of the observation that *T. vaginalis* frequently causes persistent and recurrent infections [87].

It is well recognized that the immune response to *T. vaginalis* is complex and that the outcomes of infection vary [3, 88]. The potency of inflammasome activation and differential macrophage lysis through pyroptosis by different strains may contribute to the observed differences in host responses. Our work has identified ATP sensing and $K^+$ efflux as activators of NLRP3 inflammasomes in macrophages responding to *T. vaginalis* infection. These general processes are likely to be elicited in other cell types, such as the vaginal and cervical cells first encountered by the parasite and immune cells recruited to the infection site. Our work has set the foundation for further investigation into how *T. vaginalis* activates inflammasomes and how this molecular pathway contributes to IL-1β-mediated immune responses and proinflammatory host cell death.

### Statement of Ethics

The animal work was approved by and conducted in accordance with the UCSD Institutional Animal Care and Use Committee (IACUC), protocol No. S00227M. UCSD is accredited by the American Association for Accreditation of Laboratory Animal Care (AAALAC).

### Disclosure Statement

The authors have no conflicts of interest to declare.

### Author Contributions

A.M.R. and V.N. conceived the study. A.M.R., J.A.V., K.A.P., S.D.B., X.Y.Q., and C.-M.T. conducted the experiments. A.M.R., J.A.V., K.A.P., and C.-M.T. provided key insights regarding reagents, the experimental design, and data interpretation. A.M.R. and V.N. wrote the paper, with all authors providing critical input and edits.

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