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***Chlamydia trachomatis* infection results in a modest pro-inflammatory cytokine response and a decrease in T cell chemokine secretion in human polarized endocervical epithelial cells**

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Abstract

The endocervical epithelium is a major reservoir for *Chlamydia trachomatis* in women, and genital infections are extended in their duration. Epithelial cells act as mucosal sentinels by secreting cytokines and chemokines in response to pathogen challenge and infection. We therefore determined the signature cytokine and chemokine response of primary-like endocervix-derived epithelial cells in response to a common genital serovar (D) of *C. trachomatis*. For these studies, we used a recently-established polarized, immortalized, endocervical epithelial cell model (poA2EN) that maintains, *in vitro*, the architectural and functional characteristics of endocervical epithelial cells *in vivo* including the production of pro-inflammatory cytokines. PoA2EN cells were susceptible to *C. trachomatis* infection, and chlamydiae in these cells underwent a normal developmental cycle as determined by a one-step growth curve. IL1 α protein levels were increased in both apical and basolateral secretions of *C. trachomatis* infected poA2EN cells, but this response did not occur until 72 hours after infection. Furthermore, protein levels of the pro-inflammatory cytokines and chemokines IL6, TNF α and CXCL8 were not significantly different between *C. trachomatis* infected poA2EN cells and mock infected cells at any time during the chlamydial developmental cycle up to 120 hours post-infection. Intriguingly, *C. trachomatis* infection resulted in a significant decrease in the constitutive secretion of T cell chemokines IP10 and RANTES, and this required a productive *C. trachomatis* infection. Examination of anti-inflammatory cytokines revealed a high constitutive apical secretion of IL1ra from poA2EN cells that was not significantly modulated by *C. trachomatis* infection. IL-11 was induced by *C. trachomatis*, although only from the basolateral membrane. These results suggest that *C. trachomatis* can use evasion strategies to circumvent a robust pro-inflammatory cytokine and chemokine response. These evasion strategies, together with the inherent immune repertoire of endocervical epithelial cells, may aid chlamydiae in establishing, and possibly sustaining, an intracellular niche in microenvironments of the endocervix *in vivo*.

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Keywords

Chlamydia; cytokine; endocervix; epithelial; immune; pro-inflammatory

1. Introduction

Genital serovariants (D-K) of the obligate intracellular bacterium *Chlamydia trachomatis* are the world's most common sexually transmitted bacterial pathogens, accounting for an estimated 90 million new cases reported annually [1]. *C. trachomatis* exhibits a tropism for the columnar epithelial cells of the genital mucosae, with the endocervix being the most commonly infected site in women. In a proportion of infected women, organisms also ascend into the endometrium and Fallopian tubes where chronic infection can lead to devastating reproductive consequences, including pelvic inflammatory disease (PID), tubal infertility, and ectopic pregnancy, all of which result from immune mediated damage [1]. The reason why *C. trachomatis* can cause extended infections, lasting months to years in the face of an immune response [2-6], is not well understood, but does suggest the organism can adapt to, or evade, elements of the local host immune response.

Chlamydiae have a biphasic developmental cycle that begins when non-metabolically active, infectious, elementary bodies (EBs) encounter the apical surface of polarized epithelial cells. Following entry into the host cell, EBs escape lysosomal fusion, and endosomes containing EBs fuse to form the membrane bound vacuole termed an inclusion. EBs differentiate into metabolically active, non-infectious reticulate bodies (RBs) that undergo DNA replication and binary fission. RBs then re-differentiate into EBs that may then escape the host cell through lysis or extrusion mechanisms [7, 8].

Traditional methods for culturing *C. trachomatis in vitro* utilize either murine fibroblast cell lines or the ectocervix derived cervical carcinoma cell line (HeLa). Recent studies, however, have highlighted the importance of the cell type in which chlamydiae are grown, as cell lines derived from different anatomical sites yield different growth rates and infectious yields [9, 10]. Neither HeLa cells nor murine fibroblast cells accurately represent the target cells *in vivo*, the polarized columnar epithelial cells, with respect to morphology, expression of innate immune mediators or responsiveness to TLR agonists [11, 12]. Furthermore, innate immune signaling pathways have been altered over the years during the time that HeLa cells have been grown *in vitro* [13, 14]. Alterations in innate immune pathways are significant, as it has been hypothesized that the epithelial cells, generally believed to be the only sites of chlamydial replication in the female reproductive tract (FRT), are responsible for eliciting and sustaining the inflammatory immune cascade associated with chlamydial disease via the release of intracellular IL1 α upon chlamydiae induced host cell lysis [15, 16]. Therefore, traditional cell lines may not be the optimal model systems in which to investigate the innate immune cascade elicited by *C. trachomatis* infected epithelial cells.

In recent years the orientation of the cells used to culture chlamydiae has been shown to influence chlamydial biology. Columnar epithelial cells, the target cells for chlamydial infection, maintain functionally distinct apical and basolateral membrane domains that are separated by tight junctions. Epithelial cells grown in a polarized orientation contain greater nutrient pools that are important for chlamydial growth, such as tryptophan, than their traditionally-grown submerged cell counterparts grown on plastic surfaces [17]. The use of polarized epithelial cell culture models for chlamydial studies, pioneered by Wyrick, has also revealed differences in chlamydial entry and exit mechanisms, infectious progeny, duration of the developmental cycle, infectivity, duration of the persistent growth state, reactivity to antibiotics, responsiveness to female sex steroid hormones, and innate

inflammatory responses (reviewed in [18]). All of these parameters of chlamydial biology may also influence the subsequent innate epithelial immune response to the bacteria as well, although this has not yet been investigated in more primary-like genital epithelial cells.

We recently developed an epithelial cell model derived from human endocervical tissue (A2EN cells). A2EN cells polarize and appropriately express many of the functional proteins of the endocervical epithelium *in vivo* such as hormone receptors, mucins, anti-microbial peptides, and cytokines [11]. The aim of this study was to determine the characteristics of, and the cytokine response to, *C. trachomatis* infection in polarized A2EN (polA2EN) cells in order to examine the potential role of the endocervical epithelium in initiating, sustaining and amplifying a pro-inflammatory immune response to this organism.

2. Materials and Methods

2.1 Epithelial Cell Culture

The A2EN human endocervical epithelial cell line was originally generated in our laboratory from primary epithelial cells grown out from an endocervical explant and immortalized using human papilloma virus genes E6 and E7, as recently described [11, 12]. A2EN cells were grown in a phenol red-free serum-free, medium (EpiLife; Cascade Biologics) with a defined growth supplement, and seeded onto 0.4 μ m transwell inserts under differentiation conditions to induce polarization, as previously described [11]. For comparative studies, primary endocervical epithelial cells were generated from endocervical tissue explants obtained from discarded hysterectomy tissue, as previously described [11, 12]. Collection of hysterectomy tissue was approved by the Louisiana State University Health Sciences Center Institutional Review Board. Primary and A2EN cells were seeded onto transwell membrane inserts, and polarized, as previously described [11].

HeLa 229 cells (CCL2) were obtained from the American Type Culture Collection (ATCC) and were grown in RPMI1640 supplemented with 10% fetal bovine serum.

All epithelial cells were grown and maintained in an atmosphere of 5% CO₂ at 37°C.

2.2 *Chlamydia trachomatis* Infection

PolA2EN or primary endocervical epithelial cells were infected with laboratory adapted biovariants: *C. trachomatis* serovar D (D/UW3/Cx) or serovar L2/434/Bu that were purified using Opti-Prep Density Gradient Medium (Sigma) [19] and diluted in sucrose-phosphate-glutamic acid buffer (SPG). Using the rocking method for infection, culture medium was removed from cells, and chlamydial inoculum in SPG was overlaid onto cells for 2 hours at 37°C with rocking. SPG containing chlamydial inoculum was then removed, and fresh culture medium added. For centrifugation protocols, chlamydial inoculum in SPG was placed directly into the culture medium, and cultures were centrifuged for 40 minutes at 1825xg at 35°C. Infected cells were then incubated for up to 72 hours at 37°C in culture medium. Cyclohexamide (2 μ g/mL) was added to the medium for infectivity assays only.

2.3 Inactivation of *C. trachomatis* Using Ultraviolet Light

Live *C. trachomatis* serovar D in SPG were placed in 6 well plates (1 mL of stock per well) under a laminar flow biocabinet on a rocker at a distance of 76cm from a 30 Watt ultraviolet light. Chlamydiae were rocked for 10 minutes under the UV light, sonicated using a water bath sonicator for 15 seconds to break up clumps of chlamydial particles, aliquoted, and stored at -80°C. Monolayers of HeLa 229 cells were infected with 10-fold dilutions of UV inactivated chlamydial stocks in SPG for 48 hours, fixed, and immuno-stained for chlamydial LPS to confirm that the UV-inactivated chlamydial particles were not viable and

did not form inclusions. polA2EN cells were exposed to an amount of UV inactivated chlamydiae equivalent to the amount of live bacteria required for a >95% infection rate.

2.4 *C. trachomatis* Infectivity and One Step Growth Curve Assays

Monolayers of HeLa 229 cells and polA2EN cells were infected with 0.125, 0.25, 0.5, and 1.0 μ L of *C. trachomatis* serovar D semi-purified inoculum using the protocol in section 2.2 with cyclohexamide. Infected cells were fixed with 4% paraformaldehyde for 10 minutes at room temperature followed by permeabilization with 0.1% Triton X in PBS for 10 minutes. Fixed cells were then blocked with Background Sniper (Biocare) for 30 minutes at room temperature, and stained with a FITC conjugated anti-chlamydial LPS antibody (Meridian Diagnostics) for 30 minutes at room temperature, followed by washing with PBS. Cells were then counterstained with 4',6-diamidino-2-phenylindole (DAPI) (Mol. Probes) for 1 minute at room temperature followed by washing with PBS. Membranes from polarized cultures were excised and mounted using ProLong Gold Mounting Media (Mol. Probes). Cell numbers and inclusion numbers were counted in 10 microscopic fields at 400 \times to obtain percent infectivity values. Using the results from this assay, all subsequent experiments for this manuscript were performed such that cells were infected with an amount of *C. trachomatis* inoculum that yielded a >95% infection rate to determine the direct effects of chlamydiae on the innate cytokine response in infected cells rather than infected and bystander non-infected cells.

In order to establish *C. trachomatis* growth curves, polA2EN cells were infected with *C. trachomatis* serovar D to yield a >95% infection rate. Infected cells were scraped and sonicated for 15 seconds using a water bath sonicator at 0, 12, 24, 36, 48, and 72 hours post infection. Ten-fold dilutions were performed in SPG and used to infect monolayers of HeLa 229 cells as described in section 3.3.2. Infected HeLa 229 cells were fixed 48 hours post-infection and stained with a FITC conjugated anti-chlamydial LPS antibody (Meridian Diagnostics) in order to calculate inclusion forming units (IFU), as previously described [20].

2.5 Transepithelial Electrical Resistance Measurements

A2EN cells were seeded onto transwell membrane inserts as previously described [11]. Five days post-differentiation, cells were infected with *C. trachomatis* serovar D to yield >95% infection rate. Transepithelial electrical resistance (TEER) measurements were obtained from mock-infected polA2EN cells and *C. trachomatis* infected polA2EN cells using a voltohmmeter (World Precision Instruments) every 24 hours from day 1 post-differentiation to day 9 post differentiation [11].

2.6 Microscopy and Imaging

2.6.1 Immunofluorescence Microscopy—*C. trachomatis* serovar D-infected polA2EN cells were fixed at 48 hours post-infection and stained using a FITC-conjugated anti-chlamydial LPS antibody (Meridian Diagnostics), and counterstained with DAPI as described in section 3.3.4. Fluorescent images were captured using a Leica DMRXA automated upright epifluorescence microscope (Leica Microsystems, Bannockburn, IL), a Sencicam QE chargecoupled device (Cooke Corp., Auburn Hills, MI), and filter sets optimized for Alexa 488 (exciter HQ480/20, dichroic Q495LP, and emitter HQ510/20m) and 4',6-diamidino-2-phenylindole (exciter 360/40x, dichroic 400DCLP, and emitter GG420LP). Z-axis plane capture, deconvolution, and analyses were performed with Slidebook™ deconvolution software (Intelligent Imaging Innovations, Denver, CO).

2.6.2 Transmission Electron Microscopy—For ultrastructural studies, *C. trachomatis* serovar D infected polA2EN cells were fixed in 2% paraformaldehyde/2.5% glutaraldehyde

in 200mM phosphate buffer for 1 hour at room temperature, followed by washing with PBS three times [21]. Samples were then processed and imaged for transmission electron microscopy as previously described [21].

2.7 Cytometric Bead Array Assay

HeLa 229 cells were infected with either *C. trachomatis* serovar D or LGV2 to achieve a >95% infection rate. Supernatants from infected HeLa 229 cell monolayers were collected from both mock-infected and *C. trachomatis*-infected cells at 24, 36, 48, and 72 hours post-infection and analyzed for IL6 and CXCL8 secretion using a cytometric bead assay (Millipore).

PolA2EN cells were infected with *C. trachomatis* serovars D and LGV2 to achieve a >95% infection rate. Supernatants from apical and basolateral chambers from both infected and mock-infected cells were collected over time at 24, 36, 48, and 72 hours post-infection and analyzed with a cytometric bead array that initially included (i) the pro-inflammatory cytokines Interleukin (IL)1 α , IL1 β , IL6, Tumor Necrosis Factor α (TNF α), IL17, Interferon γ (IFN γ), IL12(p70); (ii) Th2 cytokines IL4, IL5, IL7, IL9, IL10, IL15; (iii) the chemokines Eotaxin, CXCL8, Macrophage Inflammatory Protein (MIP)1 α , MIP1 β , Interferon Inducible Protein 10 (IP10), Regulated upon Activation, Normal T-cell Expressed (RANTES or CCL5), and Monocyte Chemoattractant Protein (MCP)-1, and (iv) the immunoregulatory cytokines IL11, IL13, and IL1 receptor antagonist (IL1ra). Both active and inactive forms of the regulatory cytokines transforming growth factor beta (TGF β) 1, 2, and 3 were also assayed. Inactive TGF β was examined by treating the supernatant samples with hydrochloric acid (HCl) followed by neutralization with sodium hydroxide (NaOH) as previously described [22]. Granulocyte Colony Stimulating Factor (G-CSF) and Granulocyte Macrophage Colony Stimulating Factor (GM-CSF) were also measured. All cytokines and chemokines were measured in either a BioPlex cytometric bead array kit (BioRad) or a Milliplex cytometric bead array kit (Millipore) according to the manufacturer's instructions. Only those cytokines that were detectable by cytometric bead array and are known to be secreted by epithelial cells were chosen for subsequent experiments.

Polarized and non-polarized primary endocervical epithelial cells were infected with *C. trachomatis* serovar D to achieve a >95% infection rate. Supernatants were collected from mock-infected and *C. trachomatis*-infected cells at 72 hours post-infection and analyzed with a cytometric bead array that included pro-inflammatory cytokines and chemokines: IL1 α , IL1 β , IL6, TNF α , CXCL8, MIP1 β , IP10, RANTES, G-CSF, and GM-CSF (Millipore).

2.8 Lysate and Supernatant Experiments

Mock-infected and *C. trachomatis* serovar D-infected polA2EN cells were scraped into culture medium at 24, 48, and 72 hours post-infection. Scraped cells were then sonicated for 30 seconds in a water bath sonicator to release all intracellular proteins. Lysates were then passed through a 0.2 μ m syringe filter, followed by a 0.1 μ m syringe filter to remove any chlamydial particles.

Apical and basolateral supernatants from mock-infected and *C. trachomatis* serovar D-infected polA2EN cells were also collected and filtered using the same protocol described above. Lysates and supernatants were added to non-infected polA2EN cells for 4 hours at 37°C. Lysates and apical supernatants were added to the apical chambers of polA2EN cells, whereas basolateral supernatants were also added to the basolateral chambers of polA2EN cells. Following incubation with lysates or supernatants, polA2EN cells were washed with

PBS twice, fresh culture medium was added, and then were incubated for 12 hours at 37°C. Apical and basolateral supernatants were then collected and analyzed with a cytometric bead array that included the pro-inflammatory cytokines IL1 α , IL1 β , IL6, TNF α , CXCL8, MIP1 β , IP10, RANTES, G-CSF, and GM-CSF. Apical and basolateral supernatants from untreated polA2EN cells were used as controls for baseline cytokine secretion. Lysate and supernatant experiments were designed based on previous experiments performed by Rasmussen *et al.*[15].

2.9 Statistics

For cytometric bead array assays, statistical analyses were completed using Prism software (v4.0; GraphPad, San Diego, CA). One-way analyses of variance (ANOVA) with Bonferroni posttests were employed. A P-value of <0.05 was considered significant.

3. Results

3.1 Polarized A2EN cells are susceptible to *C. trachomatis* infection

We recently developed a polarized, immortalized, endocervical epithelial cell model (polA2EN) that generates many innate immune mediators including mucin, anti-microbial peptides, cytokines and chemokines [11]. Since constitutively made innate immune mediators might interfere with *C. trachomatis* infection of polA2EN cells, the first objective of this study was to determine whether or not polA2EN cells could be infected with *C. trachomatis*. First, we utilized two methods to infect polA2EN cells with varying doses of chlamydial inoculum: the rocking method and the centrifugation method. Using the rocking method, we were only able to achieve approximately a 25% infection rate regardless of the dose used (Figure 1a) possibly because, similar to endocervical epithelial cells *in vivo*, polA2EN cells generate mucus and/or antimicrobial mediators [11], which may impede chlamydial attachment and entry. Using the centrifugation method, however, a >95% infection rate was consistently achieved (Figure 1b), although a higher volume of chlamydial inoculum was required to yield this infection rate compared to HeLa 229 cells (data not shown). Therefore, for all subsequent experiments in this study, the centrifugation method was employed to achieve a > 95% infection rate in polA2EN cells.

It has been reported that *C. trachomatis* infection can result in the degradation of epithelial junctional complex proteins, such as nectin-1 [23], which may compromise the integrity of the epithelial cell barrier. We, therefore, verified that the polarized monolayers of A2EN cells were not significantly compromised upon these conditions of chlamydial infection, as this would interfere with our investigation of differential cytokine responses from the apical and basolateral membrane compartments of polA2EN cells. We measured TEERs over a period of 4 days after chlamydial infection, and although the mean TEERs were slightly lower in *C. trachomatis* infected cells at 24-72 hours post infection compared to mock infected cells, these differences were not statistically significant (Figure 1c). TEER measurements in both mock-infected and *C. trachomatis*-infected polA2EN cells remained above 1000 Ω /cm², which is the threshold by which we measure polarization [11]. These results suggest that the polarized monolayers of A2EN cells can remain relatively intact up to 4 days after chlamydial infection.

In order to verify that chlamydiae underwent a normal developmental cycle in polA2EN cells, we employed a one-step growth curve technique using *C. trachomatis* serovar D (Figure 2a). Chlamydiae grown in polA2EN cells exhibited a latency period up to 24 hours post-infection followed by a rapid recrudescence of infectious organisms with titers peaking around 48-72 hours. Although fewer infectious EBs entered polA2EN cells compared to HeLa 229 cells, chlamydiae underwent robust replication that yielded infectious EBs equivalent to numbers of infectious EBs observed in infected HeLa cells (HeLa cell data not

shown). These results suggest that the polarized orientation of the A2EN cells creates an intracellular environment that allows for rapid productive replication of chlamydial particles, similar to that reported in polarized endometrial cells [9]. In support of the TEER data, using light microscopy, we observed minimal host cell lysis even after 72 hours in *C. trachomatis* infected polA2EN cells similar to observations made on *C. trachomatis* infected trophoblast cells [24]. *C. trachomatis* infections were also carried out to 96 and 120 hours post-infection in polA2EN cells, and we observed no evidence of significant host cell lysis unlike that observed in infected HeLa cell cultures (data not shown) [7]. These results suggest that the cell type and polarized orientation of the cells may alter the mechanism of chlamydial exit as previously hypothesized by the Wyrick group [25].

Using light microscopy, inclusions were visible by 24 hours post infection in polA2EN cells. Morphological characteristics of chlamydiae at 72 hours post-infection in polA2EN cells are shown using immunostaining against chlamydial LPS, and TEM imaging, in Figure 2b and 2c. Inclusions observed in polA2EN cells contained numerous EB at 72 hours post-infection (Figure 2b, c) and were consistent with healthy chlamydial growth.

3.2 *C. trachomatis* does not initiate a robust inflammatory cytokine/chemokine response in polA2EN cells

HeLa cells infected with *C. trachomatis* LGV2 robustly up-regulate the secretion of pro-inflammatory cytokines, including IL6, CXCL8, GRO α , GM-CSF, and IL1 α [15, 26, 27]. We, first, confirmed these results in HeLa 229 cells using both *C. trachomatis* serovar D as well as LGV 2. Similar to results found in previous studies [15, 26, 28], we observed a significant up-regulation of both CXCL8 and IL6 by 48 hours following *C. trachomatis* infection with either serovariant (data not shown). Although we observed a significant up-regulation of pro-inflammatory cytokines in infected HeLa 229 cells, we did not observe the magnitude of up-regulation of pro-inflammatory cytokines (data not shown), particularly CXCL8 (~23 fold), that Rasmussen *et al.* observed using *C. trachomatis* serovar D [15]. Our results were more consistent with results observed by Dessus-Babus *et al.* who observed a more modest up-regulation of CXCL8 and IL6 upon chlamydial infection of HeLa 229 cells [28].

Next, the pro-inflammatory cytokine profiles in *C. trachomatis* serovar D infected polA2EN cells were determined. PolA2EN cells were infected with *C. trachomatis* serovar D to yield an infection rate of >95%. Apical and basolateral supernatants from *C. trachomatis* infected polA2EN cells were collected and subjected to cytometric bead array analyses. We initially assayed for an extended panel of cytokine analytes, which included IL1 α , IL1 β , IL2, IL4, IL5, IL6, IL7, CXCL8, IL9, IL12(p70), IL13, IL15, IL17, GM-CSF, G-CSF, TNF α , IFN γ , MIP1 α , MIP1 β , IP10, RANTES, MCP-1, and Eotaxin. Only those cytokines that were secreted by polA2EN cells were chosen for further studies. As described previously, polA2EN cells, similar to other primary-like epithelial cells [29, 30], constitutively secreted CXCL8, IL6, IP10, and RANTES (Figure 3a,b,e,f) as well as G-CSF and GM-CSF (data not shown) [11]. Unlike the results observed in *C. trachomatis* infected HeLa cells by Rasmussen *et al.*, we observed no significant up-regulation of IL6, CXCL8 or TNF α at any timepoint over the course of the developmental cycle and up to 120 hours post-infection (Figure 3a-c; Supplemental Figure 1a-c). Similar to infected HeLa cells [15], we observed a modest but significant increase in IL1 α secretion (Figure 3d, Supplemental Figure 1c) (approximately 4 fold; $p < 0.001$), and this did not occur until 72 hours post-infection. Intriguingly, we consistently found a specific, and highly significant, decrease in the constitutive secretion of IP10 and RANTES by 72 hours post-infection that lasted up to 120 hours post-infection. The latter results suggest that *C. trachomatis* may actively modulate these T cell chemokines (Figure 3e,f; Supplemental Figure 1e,f; $p < 0.001$). We did not observe a significant modulation in secretion of any other analytes measured upon *C.*

trachomatis infection even when the inoculum dose was increased up to 4 times the amount necessary to achieve >95% infection (representative CXCL8 graph shown in Supplemental Figure 2).

We next investigated whether polarization of the A2EN cells contributed to the lack of pro-inflammatory cytokine secretion upon *C. trachomatis* infection. We, therefore, infected A2EN cells grown submerged in traditional plastic culture plates at an infection rate of >95%. Similar to results observed in *C. trachomatis* infected polA2EN cells, infection of non-polarized A2EN cells failed to result in a significant up-regulation in IL6, CXCL8 or TNF α (Figure 4a,b,c). Also similar to results observed in *C. trachomatis* infected polA2EN cells, IP10 and RANTES were decreased in *C. trachomatis* infected non-polarized A2EN cells (Figure 4d,e). These results suggest that the polarized orientation of A2EN cells does not alter the pro-inflammatory immune response.

Because no significant pro-inflammatory cytokine response was observed in *C. trachomatis* serovar D infected polA2EN cells, we determined whether cytokines involved in anti-inflammatory processes might be modulated upon infection. We analyzed TGF β 1, 2, and 3 as well as IL1 α , IL11, IL13, and IL10 secretion upon *C. trachomatis* infection. PolA2EN cells did not secrete TGF β 1 or 3 (data not shown). PolA2EN cells also did not secrete detectable levels of IL13, or IL10 (data not shown). TGF β 2 was detectable in polA2EN cell secretions, however it was unaltered upon *C. trachomatis* infection (Figure 5a). PolA2EN cells secreted high constitutive levels of IL1 α at their apical surface, however secretion was unaltered upon *C. trachomatis* infection (Figure 5b). Interestingly, the pleiotropic, anti-inflammatory cytokine IL11 was significantly up-regulated only in the basolateral secretions from *C. trachomatis* infected polA2EN cells (Figure 5c, p<0.001) and was undetectable in apical secretions.

We also determined whether the modest pro-inflammatory cytokine secretion in response to *C. trachomatis* was serovar specific as disseminating biovariants (LGV) and non-disseminating biovariants (D-K) have been shown to elicit different innate immune responses [15, 28]. Similar to what we observed using serovar D, polA2EN cells did not up-regulate IL6 or CXCL8 secretion upon LGV2 infection (data not shown). We also observed a decrease in IP10 and RANTES secretion upon *C. trachomatis* LGV2 infection (data not shown).

3.3 *C. trachomatis* infected primary endocervical epithelial cells have a similar cytokine profile to infected polA2EN cells

We next investigated whether the modest pro-inflammatory cytokine secretion and the decrease in IP10 and RANTES were an artifact of using an immortalized endocervical epithelial cell model. We therefore utilized polarized primary endocervical epithelial cells and infected them with *C. trachomatis* serovar D at >95% infection rate. Polarized primary endocervical epithelial cells did not significantly up-regulate the pro-inflammatory cytokines and chemokines CXCL8, IL6, and TNF α in response to *C. trachomatis* in either apical or basolateral secretions (Figure 6a-f). IL1 α secretion was increased in some patients' cells, while levels in other cells remained unchanged upon *C. trachomatis* infection (Figure 6g,h). Similar to results observed in polA2EN cells, we also observed a decrease in the T cell chemokines IP10 and RANTES secretion in polarized primary endocervical epithelial cells infected with *C. trachomatis* (Figure 7a-d). The decrease in IP10 and RANTES secretion we observed in polarized primary endocervical epithelial cells support and validate the results generated in polA2EN cells.

3.4 UV inactivated *C. trachomatis* does not elicit an IL-1 α response, nor significantly decreases the IP10 and RANTES response in polA2EN cells

We next investigated whether the modest pro-inflammatory cytokine response and the reduction in T cell chemokine secretion required productive chlamydial infection. We inactivated chlamydiae using ultraviolet light which inhibits active growth of the bacteria, while maintaining the structural integrity of molecules or pattern associated molecular patterns (PAMPs) on the bacteria's surface that may interact with pattern recognition receptors (PRRs). Similar to experiments performed with live, active bacteria, UV inactivated chlamydiae did not elicit a robust CXCL8, IL6, or TNF α response (Figure 8a-c). Interestingly, UV inactivated chlamydiae also did not increase IL1 α secretion in polA2EN cells (Figure 8d), suggesting that a productive infection is needed in order to elicit this response similar to what has been previously reported [15, 31]. UV inactivated chlamydiae also did not abrogate IP10 and RANTES secretion in polA2EN cells (Figure 8e,f), also suggesting that a productive chlamydial infection is necessary for the bacteria to cause the decrease in T cell chemokine secretion in polA2EN cells.

3.5 Neither lysates nor supernatants from *C. trachomatis* infected polA2EN cells stimulate non-infected polA2EN cells to produce pro-inflammatory cytokines

The previously reported robust secretion of CXCL8 following *C. trachomatis* infection in HeLa cells observed by Rasmussen et al. was suggested to be due to IL1 α stimulation of bystander non-infected epithelial cells resulting in the amplification of the CXCL8 response [15]. Therefore, we first investigated whether polA2EN cells could respond to IL1 α , as a lack of response to IL1 α is a possible explanation for why polarized A2EN cells failed to up-regulate pro-inflammatory cytokines upon chlamydial infection. Figure 9 shows that, upon addition of recombinant IL1 α , polarized A2EN cells significantly up-regulated the secretion of IL6, CXCL8, TNF α , IP10, MIP1 β , RANTES, GM-CSF, and G-CSF in both apical and basolateral secretions. These results indicate that non-infected polA2EN cells can respond to IL1 α to secrete pro-inflammatory cytokines.

Amplification of the pro-inflammatory cytokine response by *C. trachomatis* infected HeLa cell cultures was demonstrated to be due to IL1 α , which was assumed to be released upon host cell lysis [15]. While we also established that IL1 α secretion was significantly increased in *C. trachomatis* infected polA2EN cells, we did not observe significant epithelial cell lysis up to 120 hours post-infection. We therefore investigated whether lack of cell lysis was responsible for the absence of pro-inflammatory cytokine amplification in infected polA2EN cells. We harvested *C. trachomatis* infected polA2EN cells at 24, 48, and 72 hours post-infection, sonicated and filtered them to remove chlamydial particles. We then added the filtered lysates to the apical surfaces of non-infected polA2EN cells for 4 hours. Following incubation, cells were washed, and fresh medium was added to both the apical and basolateral chambers for 12 hours. Apical and basolateral secretions were collected and assayed for cytokines and chemokines. The secretion of IL6 and CXCL8 was unaltered between unstimulated, mock-infected lysate and infected lysate treatment conditions (72 hour timepoint shown; 24 and 48 hour timepoints not shown; Figure 10a,b). These results suggest that intracellular components, such as IL1 α , released upon host cell lysis during chlamydial infection are unable amplify the pro-inflammatory cytokine response in polA2EN cells. The secretion of IP10 and RANTES was also unaltered between unstimulated, mock-infected lysate and infected lysate treatment conditions (72 hour timepoint shown; 24 and 48 hour timepoints not shown; Figure 10c,d), suggesting that host cell lysis and the components released by infected cells do not down-modulate these T cell chemokines. There were no statistically significant differences between experimental groups for all other analytes measured.

Finally, we determined if apical and basolateral secretions from *C. trachomatis* infected polA2EN cells could stimulate non-infected cells to modulate cytokine and chemokine secretion. We collected supernatants from *C. trachomatis* infected polA2EN cells, filtered them to remove chlamydial particles, and added the filtered supernatants to non-infected polA2EN cells for 4 hours. We added apical secretions to apical surfaces, since apical membrane surfaces are likely to be exposed to apical secretions from neighboring cells *in vivo*. Likewise, we treated basolateral surfaces of polA2EN cells with the collected basolateral supernatants. Similar to the lysate protocol, we removed the supernatants, washed, and incubated the cells for 12 hours with fresh medium. Similar to results observed using *C. trachomatis* infected cell lysates, there were no statistically significant differences between the experimental groups for IL6, CXCL8, IP10 or RANTES (Figure 11a-d). There were no statistically significant differences for all other analytes measured as well.

3.5 Discussion

In order to investigate the endocervical epithelial cytokine response to *C. trachomatis*, we used a polarized, endocervical epithelial cell model that we could apically challenge, successfully infect, and subsequently differentiate apical and basolateral cytokine responses. We observed that *C. trachomatis* infection of polA2EN cells significantly increased IL1 α in both apical and basolateral secretions, but, unlike other intracellular bacteria that induce a rapid and transient IL1 α response [32], the response we observed did not occur until 72 hours post infection. We also found that a >95% *C. trachomatis* infection of polA2EN cells: (1) failed to up-regulate the secretion of the major pro-inflammatory cytokines CXCL8, IL6 or TNF α , in either apical or basolateral secretions at any time measured; (2) induced secretion of IL11, an anti-inflammatory, tissue repair cytokine, at the basolateral surface of polA2EN cells; and (3) significantly decreased the constitutive apical and basolateral secretion of IP-10 and RANTES in polA2EN cells. Overall, these observations suggest *C. trachomatis* can evade specific endocervical epithelial cytokine responses, and that these cells could provide a hospitable intracellular niche for genital *C. trachomatis* infection.

It was not entirely surprising that *C. trachomatis* failed to robustly up-regulate pro-inflammatory cytokines in polA2EN as numerous studies have documented that this bacteria uses evasion mechanisms to interfere with host immune responses [33-39]. Some of these studies have described chlamydial proteases that are secreted into the host cell cytosol they can manipulate host proteins and signaling pathways [39, 40]. For example, the NF κ B pathway may be modulated by several different chlamydial proteins and mechanisms, all of which can interfere with NF κ B-mediated gene transcription and regulation [36, 37, 41]. *C. trachomatis* interference with cytokine signaling pathways may be one explanation as to why chlamydiae failed to robustly upregulate a pro-inflammatory innate response in polA2EN cells, as many of the pro-inflammatory cytokines, such as CXCL8, IL6, TNF α and IL1, are downstream of NF κ B activation [42]. Since *C. trachomatis* infection of polA2EN cells did not cause a decrease in the constitutive secretion of the pro-inflammatory cytokines CXCL8 and IL6, there may be other innate immune signaling pathways involved in the secretion of these cytokines, such as the MAP kinase pathway [26], that are unaltered during infection. It is also unlikely that chlamydial LPS is playing a major role in the endocervical epithelial cytokine responses, since, TNF α was not upregulated in response to chlamydial exposure unlike what was reported by Ingalls et al. who found modest cytokine responses in CHO cells to be mediated by chlamydial endotoxin [43]. Neither polA2EN cells nor primary endocervical epithelial cells express high levels of the LPS receptors TLR4 or CD14 [11, 12], further suggesting that chlamydial LPS may play a minimal role in eliciting an epithelial cytokine response in the endocervix.

Previous studies have demonstrated that the robust CXCL8 response by *C. trachomatis*-infected HeLa cell cultures was primarily mediated by IL1 α [15, 44]. Although, we observed an increase in IL1 α protein levels in *C. trachomatis* infected polA2EN cells at 72 hours post infection, we did not observe an increase in CXCL8 at any point during the chlamydial developmental cycle and up to 120 hours postinfection. IL1 α release upon host cell lysis and stimulation of non-infected bystander epithelial cells was the major mechanism by which the HeLa pro-inflammatory cytokine response was amplified [15]. polA2EN cells infected with *C. trachomatis*, did not, however, exhibit significant lysis up to 120 hours post-infection. IL1 α may be an indicator of cell damage in the absence of cell lysis, however, which may be caused by chlamydiae infection [45]. It is also possible that chlamydial particles exit polA2EN cells by extrusion thus explaining the minimal cell lysis we observed [7]. We, therefore, lysed infected polA2EN cells and investigated whether these lysates, or supernatants could stimulate non-infected polA2EN cells and amplify the CXCL8 response similar to that observed in infected HeLa cells [15]. Neither lysates nor supernatants harvested from *C. trachomatis* infected polA2EN cells, however, stimulated a pro-inflammatory response in non-infected cells. These results suggest that neither intracellular nor released IL1 α , nor any other intracellular factor could amplify the CXCL8 response in non-infected polA2EN cells. Importantly, polA2EN cells and primary endocervical epithelial cells do constitutively secrete a high level of IL1ra from their apical surfaces which HeLa cells do not (data not shown for primary or HeLa cells). IL1ra inhibits the binding of IL1 α to its receptor and prevents the subsequent pro-inflammatory immune cascade [46]. The constitutive apical secretion of IL1ra, therefore, may explain why the IL1 α that is secreted during *C. trachomatis* infection is unable to stimulate non-infected cells to amplify the pro-inflammatory cytokine response. PolA2EN cells were capable of responding to exogenous IL1 α stimulation, however, it required 20ng/mL, a very high concentration of IL1 α , to induce a cytokine response. These results indicate that the high constitutive IL1ra might compete for the IL1 receptor and prevent lower concentrations of IL1 α from amplifying the pro-inflammatory cytokine response. Constitutive secretion of IL1ra has been demonstrated in other mucosal cell types and is believed to be an endogenous regulatory mechanism by which unwanted inflammation is controlled in various tissue types [47], and has been negatively associated with pathology in trachoma patients [48-50]. The fact that polA2EN cells constitutively secrete a higher level of IL1ra from the apical membrane compared to the basolateral membrane, suggests that the two membrane compartments could play distinct roles in regulating inflammatory responses. Hence, IL1ra may be constitutively secreted at high concentrations at the apical surface of the endocervical epithelium in order to prevent IL1 α from excessively amplifying the epithelial cytokine response and compromising the integrity of this barrier. Since IL1ra is not constitutively secreted, nor upregulated by chlamydial infection at the basolateral membrane, the basal increase in IL1 α secretion by infected endocervical epithelial cells could potentially still signal to underlying resident IL-1 Receptor-expressing leukocytes. It is also possible that a sustained secretion of IL1 α from the basolateral membrane compartment of infected epithelial cells could contribute to chronic inflammation that may ultimately lead to pathology in some individuals. Our results indicate that further studies on the regulation of IL1 α , IL1ra, and IL1 receptor expression are needed to determine how these factors could contribute to the establishment, clearance, chronicity and immunopathology of *C. trachomatis* infections and also other sexually transmitted diseases at this important site.

The observation that polA2EN cells infected with *C. trachomatis* fail to elicit a robust epithelial pro-inflammatory cytokine response contrasts with observations made in the murine *in vitro* model of chlamydial infection using *Chlamydia muridarum* infection in oviduct-derived epithelial cells [51, 52]. One reason for this discrepancy may be that *C. muridarum* infection fails to interfere with murine host cell signaling pathways needed to

initiate the inflammatory cascade. Christian *et al.* and others have reported that there are differences between *C. muridarum* and *C. trachomatis* with respect to how these species interfere with proteins in innate immune signaling pathways in their respective host cells [36, 41, 53]. Another possible reason for the discrepancy is that the epithelial cells are derived from two distinct anatomical sites, the endocervix and the oviduct, respectively. Epithelial cells in the oviduct, which is a relatively sterile site, may respond more robustly to organisms than cells in the endocervix which routinely encounter microflora from the lower FRT [54, 55]. The hypothesis that there are site specific immune responses in the FRT is indirectly supported by the work of Hvid *et al* who demonstrated that IL-1 is central to human fallopian tube damage during *C. trachomatis* infection. This IL1-mediated damage can be inhibited by exogenous IL-1ra, suggesting that this regulatory cytokine is not constitutively made, or upregulated to sufficient concentrations, in the Fallopian tubes [56]. Trophoblast cells infected with *C. trachomatis* also increases IL1 β via Nod1 signaling pathways [24, 57]. Taken together, these studies suggest there may be major differences in *Chlamydia's* interaction with its target host cells dependent on the species and tissue site. Understanding species and site specific interactions with chlamydial organisms is imperative as we seek to determine the reservoirs of bacteria and mechanisms by which *C. trachomatis* establishes chronic infection and induces pathology.

In this study, we also demonstrated that *C. trachomatis* infection up-regulated IL11 in basolateral secretions of polA2EN cells. These results are consistent with observations made by others in HEC-1B endometrial epithelial cells and HeLa cells [28, 58]. IL11 has also been found in ocular secretions from patients who have chronic scarring from trachoma [48]. IL11 is a pleiotropic cytokine that has multiple functions including inhibition of NF κ B activation, cytoprotection at mucosal sites, modulation of macrophage phenotypes, promotion of cell survival and differentiation, promotion of tissue repair, and reduction of inflammatory responses ([59]; reviewed in [60, 61]). The role of IL11 in tissue repair is to up-regulate collagen and other connective tissue proteins leading to fibrosis [60]. Our results suggest that the infected endocervical epithelial cells may modulate inflammatory processes associated with chlamydial disease by secreting IL11. The observation that IL11 was only increased in basolateral secretions in *C. trachomatis* infected polA2EN cells highlights the possibility that the two distinct membrane domains in polarized epithelial cells play different roles in the induction of innate immune responses and tissue repair.

The constitutive secretion of two key T lymphocyte chemokines, IP10 and RANTES, was significantly decreased in secretions from *C. trachomatis* infected polA2EN cells. IP10 and RANTES are chemokines that bind to CXCR3 and CCR5, respectively, resulting in the recruitment of CXCR3 positive and CCR5 positive leukocytes to the target tissues, which include T cells and NK cells [62, 63]. These two cell types are the major producers of interferon gamma (IFN γ), the key cytokine that can mediate resolution of *C. trachomatis* infections when infected cells are exposed to sufficiently high concentrations [64-67]. If *C. trachomatis* can subvert the epithelial cell-mediated secretion of IP10 and RANTES *in vivo*, this may prevent directed migration of IFN γ secreting T lymphocytes and NK cells into the immediate microenvironment of infected epithelial cells, particularly in pseudoglands of the endocervix [64, 65, 68, 69]. This may be significant as cell-to-cell contact between T cells and epithelial cells is required for T cells to facilitate IFN γ mediated killing of *C. trachomatis* [70]. The decrease in IP10 and RANTES upon chlamydial infection of endocervical epithelial cells suggests that type I interferon or interferon-stimulated gene (ISG) signaling pathways may also be modulated during *C. trachomatis* infection. Type I interferons and components of type I interferon signaling pathways were shown to be involved in inflammation and pathology in *in vitro* and *in vivo* murine models of *Chlamydia* [51, 71] as well as *in vitro* human airway epithelial cell models of *C. pneumoniae* [72]. Further investigation is needed to determine if *C. trachomatis* might be interfering with type

I interferons, ISGs or other proteins involved in signaling pathways that lead to the downstream transcription, translation, and secretion of IP10 and RANTES, and the functional significance in human endocervical infections.

Overall, we propose that when *C. trachomatis* survives the innate physical and secreted immune mediators in the lower FRT and endocervix, the endocervical epithelium could provide a hospitable niche for *C. trachomatis* to establish infection for a number of reasons. First, the delay in IL1 α up-regulation combined with a lack in amplification of CXCL8, IL6 and TNF α will likely delay recruitment to, and activation of, leukocytes in an infected endocervix until well after bacteria have established an infection. Second, constitutive, apically-secreted IL1ra combined with minimal host cell lysis may help protect the integrity of the epithelial cell layer by preventing or regulating the IL1-mediated amplification of a pro-inflammatory response by bystander uninfected cells. Third, IL11 may modulate the phenotype or response of infiltrating leukocytes to reduce a pro-inflammatory response. Finally, the minimal epithelial lysis during productive infection suggests that the bacteria may be able to maintain an intracellular niche in some of the cells it originally infects in addition to infecting new cells. These factors, in combination with the relatively slow *in vivo* turnover of human endocervical epithelial cells [73], the ability of *C. trachomatis* to extend the life of infected cells by modulating apoptosis-inducing pathways [74-77], the evasion of IFN γ mediated mechanisms of clearance [20, 53, 78-80], and/or the evasion of key effectors of acquired immunity [34, 35, 81], and the evasion of IFN γ mediated mechanisms of clearance [34, 38], may explain why some infections take significantly extended times to clear. Taken together, these complex factors suggest that foci of infected cells could be maintained in the endocervix even as others are cleared. Though chlamydiae may utilize evasion strategies to circumvent or alter the immune response, our results do not rule out the possibility that infections of endocervical epithelial cells may also induce a significant inflammatory response, as sustained IL α secretion at the basolateral membrane of infected endocervical epithelial cells may overcome evasion mechanisms utilized by chlamydiae and subsequently activate underlying macrophages or dendritic cells. These immune cells are not generally believed to be permissive for productive genital chlamydial infection [82, 83], and therefore may not be subject to active subversive strategies used by the bacteria. Hence, they may activate and amplify pro-inflammatory innate and adaptive immune responses. Whether an ensuing leukocytic infiltrate could clear infection and/or mediate damage and how long this takes is likely dependent upon multiple factors that influence the endocervical micromilieu which is critical for understanding how to prevent chlamydial disease. Furthermore, it is important to investigate the dynamics between the infected host cells and chlamydiae and the ensuing epithelial innate immune response, as long-term survival of bacteria may likely require them to enter into alternative growth forms, which are believed to contribute to chlamydial disease and are more refractory to antibiotic treatment [84, 85].

There are many other confounding factors that influence and likely determine whether an endocervical chlamydial infection clears, sustains, ascends into the endometrium, and/or mediates damage. One may be the initial dose of chlamydial inoculum. If there are high numbers of infectious chlamydial particles in the inoculum, a greater proportion of endocervical epithelial cells will be infected; this may result in an increased concentration of IL1 α at the basolateral surface. IL1 α , in the absence of an increased IL1ra, may elicit a pro-inflammatory immune response that then may compromise the epithelial barrier. Higher numbers of infectious particles and an increase in IL1 α may also change the ratio of the infected cells to non-infected bystander cells, and hence, alter the immune and nutrient milieu in this locale [34, 81, 86]. It is also more likely that upper FRT tissues could be exposed to bacterial particles and become infected when there is a higher initial infectious inoculum. The ratio of dead bacteria to live bacteria may also influence the outcome of

chlamydial infection of the endocervix since replicating and non-replicating chlamydiae induce different IL1 α and T cell chemokine responses. The presence of a pre-existing inflammatory infiltrate, often observed in normal transformation zone and endocervix tissues, or associated with a coinfection, may also influence the early dynamics, and hence outcome, of infection via the immediate exposure of these leukocytes to bacteria and IL1 α [87, 88]. Other co-factors include the presence of female steroid hormones and/or synthetic progestins that may influence acquisition of chlamydial infection [89-92]; relative exposure to TGF β -rich, immunomodulatory, seminal plasma [93, 94], and vaginal co-infections such as bacterial vaginosis (BV) and trichomoniasis, both of which may provide a source of indole in a tryptophan-depleted microenvironment and therefore aid in the maintenance of a bacterial reservoir in the presence of IFN γ in the endocervix [53, 78-80, 95].

In summary, the data generated in this study provide valuable information on the innate cytokine repertoire of human endocervical epithelial cells and the evasion strategies used by the *C. trachomatis* to modify some of these innate responses. *In vivo* exogenous cofactors will also influence this relationship. As we move forward, the polA2EN model will be valuable in establishing the role of, and mechanism by which, these factors influence bacterial growth, survival, and subsequent modulation of the innate immune response in the endocervical epithelium. Generation of similar endometrial and Fallopian tube epithelial cell models will also allow us to examine site specific differences in these responses. Together, these studies may enlighten us to why and how human *C. trachomatis* infections can be so extended.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

polA2EN	polarized endocervical epithelial cells
SPG	sucrose phosphate glutamic acid buffer
IFU	inclusion forming units
FRT	female reproductive tract
PID	pelvic inflammatory disease
EBs	elementary bodies

RBs	reticulate bodies
DAPI	4', 6-diamidino-2-phenylindole
TEER	transepithelial electrical resistance

Highlights for Buckner *et al.* **Cytokine**

We infected a novel polarized endocervical epithelial model with *C. trachomatis*

C. trachomatis infection failed to yield a robust pro-inflammatory cytokine response

Constitutive RANTES and IP10 were decreased by *C. trachomatis* infection

C. trachomatis infected primary cells yielded similar cytokine responses Results suggest

C. trachomatis uses evasive strategies to survive in the endocervix

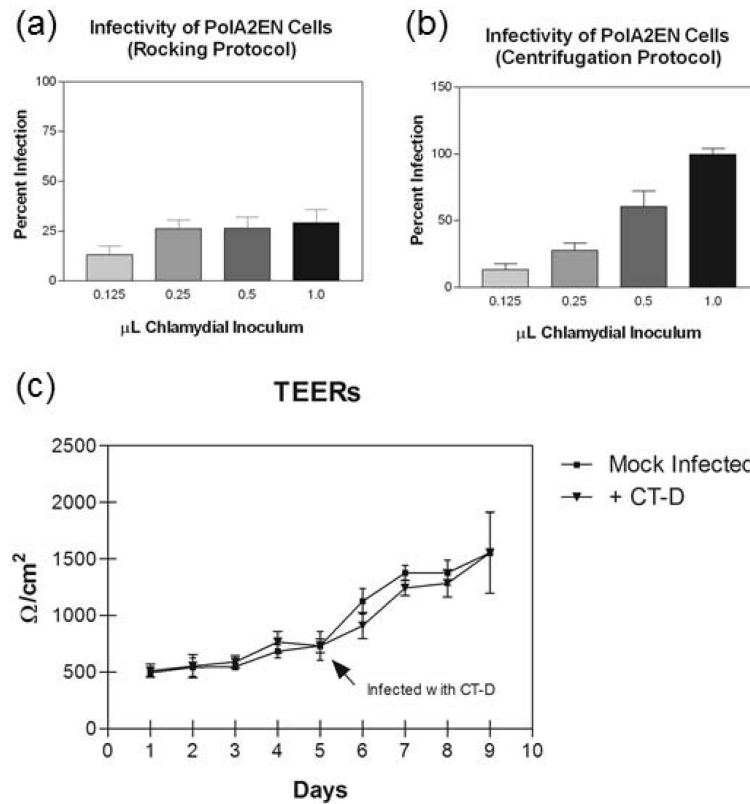


Figure 1. PolA2EN cells are susceptible to *C. trachomatis* serovar D infection, and chlamydial infection does not significantly compromise the polarized monolayer

(a) Using the rocking protocol, infection rates of *C. trachomatis* in polA2EN cells never exceeded approximately 25% no matter the dose of chlamydial inoculum used. (b) Using the centrifugation protocol for *C. trachomatis* infection, polA2EN cells were capable of being infected at an infection rate of >95% when 1.0 μL chlamydial inoculum was used. No significant cell lysis was observed in infected polA2EN cells. Infection rates shown are mean levels \pm the standard deviation (SD) from 3 independent experiments each performed in triplicate. (c) TEER values were not statistically significant between mock infected cells and *C. trachomatis* infected cells up to 4 days post-infection. Data points represent mean values \pm SD pooled from 3 independent experiments each performed in triplicate.

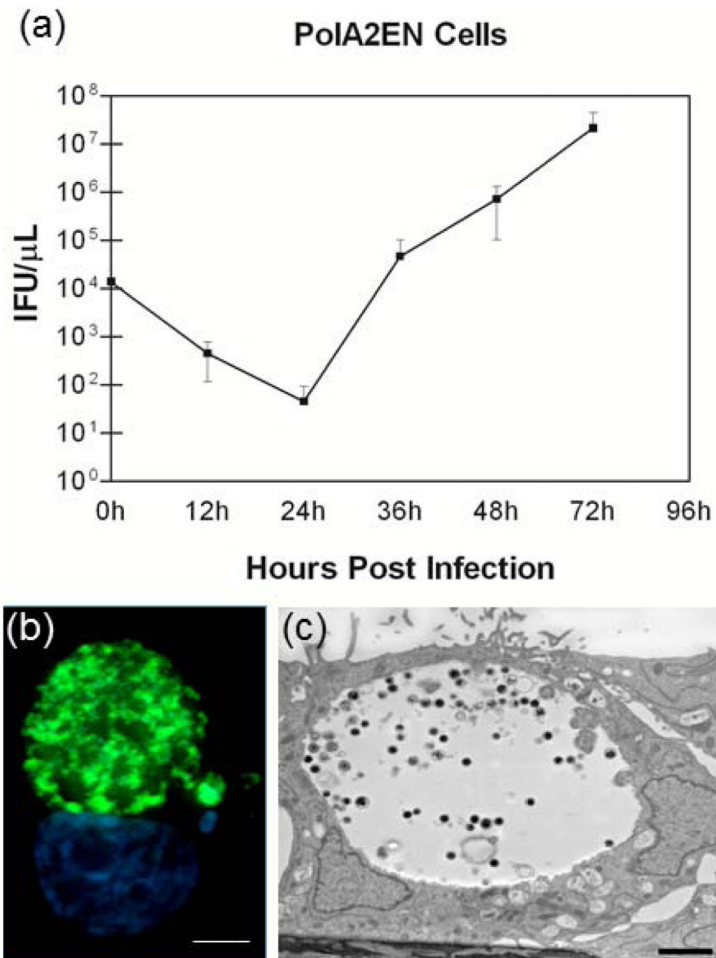


Figure 2. *C. trachomatis* in polA2EN cells exhibits a normal growth pattern and morphology
 (a) IFUs at time 0h post-infection represent chlamydiae that have entered the cells after centrifugation. Values represent mean values \pm SD pooled from 3 independent experiments each performed in triplicate. (b) PolA2EN cells were infected with *C. trachomatis* serovar D, fixed at 72 hours post-infection and immunostained with a FITC conjugated α -LPS antibody (green) and counterstained with DAPI (blue) (1000x magnification). Scale bar represents 5 μ m. (c) *C. trachomatis* infected polA2EN cells were fixed at 72 hours post-infection and processed for TEM. Dense particles representing EBs were visible. Scale bar represents 1 μ m.

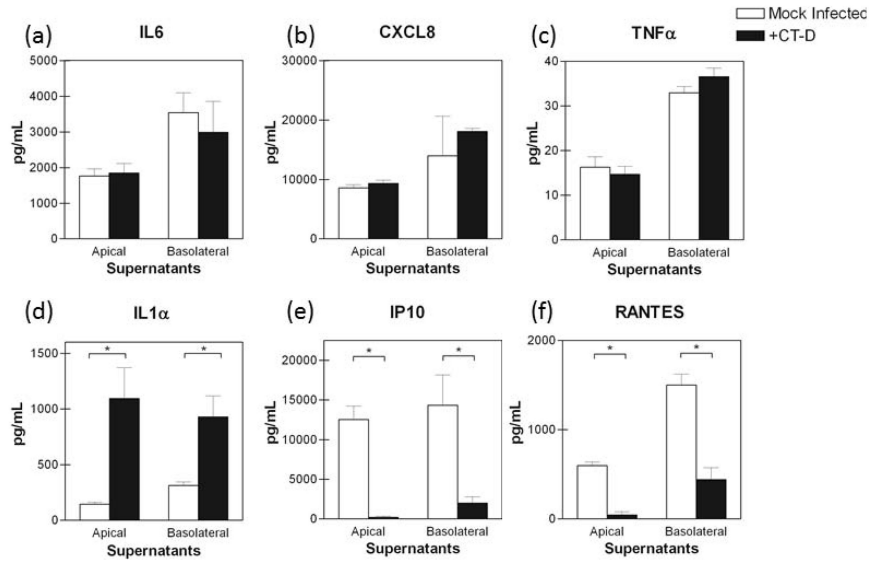


Figure 3. *C. trachomatis* fails to elicit a robust pro-inflammatory cytokine response and decreases T cell chemokine secretion in polA2EN cells

PolA2EN cells were infected with *C. trachomatis* serovar D (>95% infection rate). Apical and basolateral supernatants were subjected to cytometric bead array analyses for (a) IL6, (b) CXCL8, (c) TNFα, (d) IL1α, (e) IP10, and (f) RANTES. Graphs shown represent the 72 hours post-infection; cytokine values represent mean values ± SD. Graphs shown are representative graphs from 3 independent experiments each performed in triplicate. Statistical analyses between mock infected and *C. trachomatis* infected parameters were performed using a One-way ANOVA (*p<0.001).

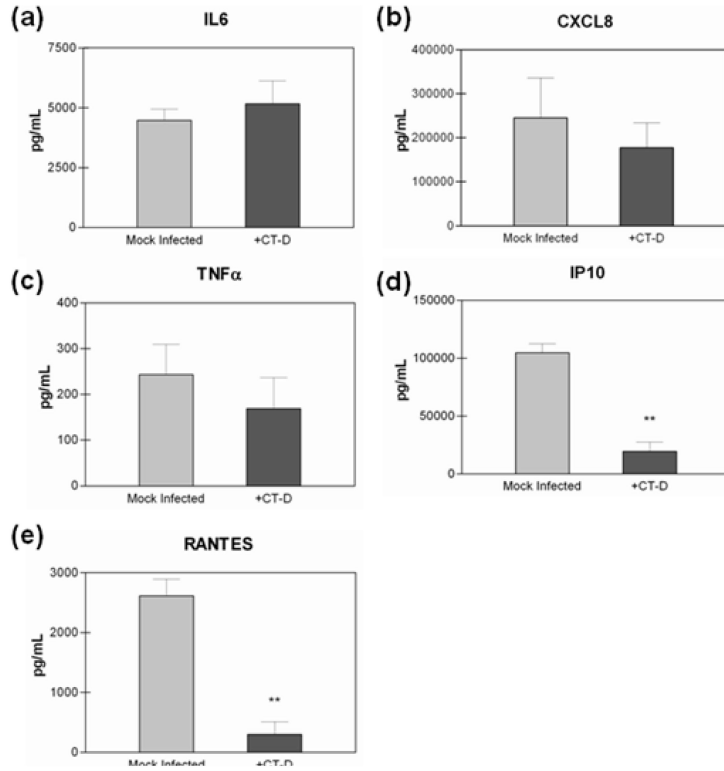


Figure 4. *C. trachomatis* fails to elicit a pro-inflammatory cytokine response, and decreases T cell chemokines in A2EN cells grown in submerged culture

A2EN cells grown in submerged culture were infected with *C. trachomatis* serovar D (>95% infection rate). Supernatants were collected at 72 hours post-infection and subjected to cytometric bead analyses for (a) IL6, (b) CXCL8, (c) TNF α , (d) IP10, and (e) RANTES. Cytokine measurements represent mean values \pm SD. Graphs shown are representative graphs from 2 independent experiments each performed in triplicate. Statistical comparisons were performed between mock infected cell and infected cell parameters using a one-way ANOVA (** $p < 0.001$).

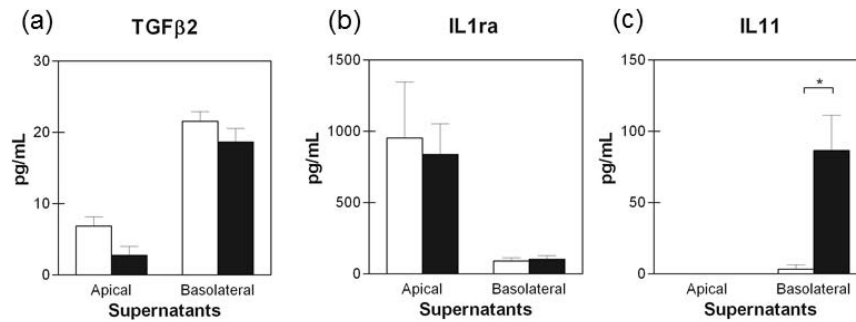


Figure 5. *C. trachomatis* elicits an IL11 response, but not TGFβ or IL1ra in polA2EN cells
 PolA2EN cells were infected with *C. trachomatis* serovar D (>95% infection rate). Apical and basolateral supernatants were collected at 72 hours post-infection and subjected to cytometric bead array analyses for (a) TGFβ2, (b) IL1ra, and (c) IL11. Cytokine values represent mean values \pm SD pooled from 3 independent experiments each performed in triplicate. Statistical analyses between mock infected and *C. trachomatis* infected parameters were performed using a One-way ANOVA (* $p < 0.001$).

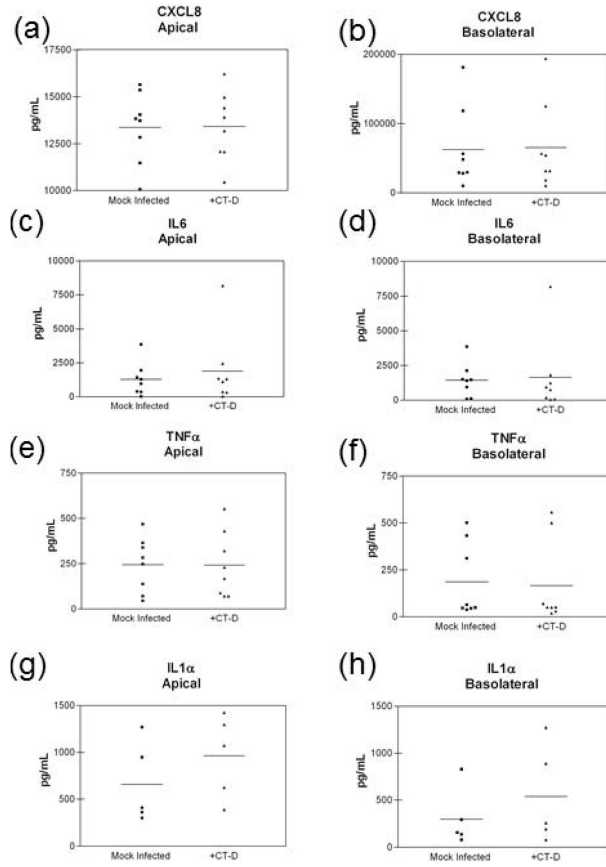


Figure 6. *C. trachomatis* infected primary endocervical epithelial cells exhibit a similar cytokine pattern to infected polA2EN cells

Primary endocervical epithelial cells (n=5-8) were infected with *C. trachomatis* serovar D (>95% infection rate). Apical and basolateral supernatants were collected at 72 hours post-infection and subjected to cytometric bead array analyses for (a,b) CXCL8, (c,d) IL6, (e,f) TNF α , and (g,h) IL1 α . Cytokine values represent mean values. Each experiment was performed in triplicate. Statistical analyses between mock infected and *C. trachomatis* infected parameters were performed using a Wilcoxon nonparametric matched pairs test.

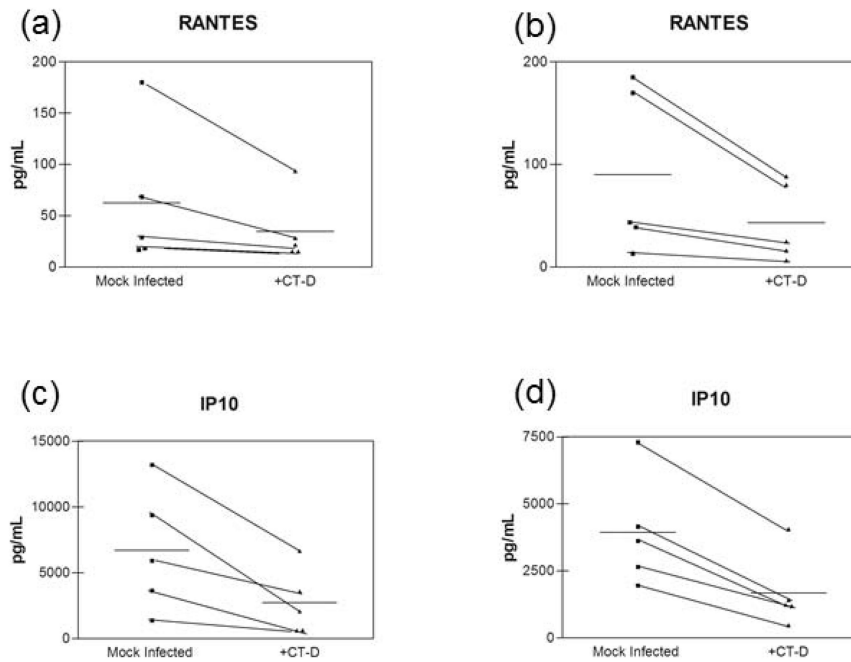


Figure 7. *C. trachomatis* infected primary endocervical epithelial cells exhibit a decrease in IP10 and RANTES

Primary endocervical epithelial cells (n=5) were infected with *C. trachomatis* serovar D (>95% infection rate). Apical and basolateral supernatants were collected at 72 hours post-infection and subjected to cytometric bead array analyses for (a,b) IP10, and (c,d) RANTES. Cytokine values represent mean values. Overall mean indicated by horizontal line. Each experiment was performed in triplicate. Statistical analyses between mock infected and *C. trachomatis* infected parameters were performed using a Wilcoxon nonparametric matched pairs test. Statistical analyses revealed that the decrease in both IP10 and RANTES was approaching significance ($p < 0.06$).

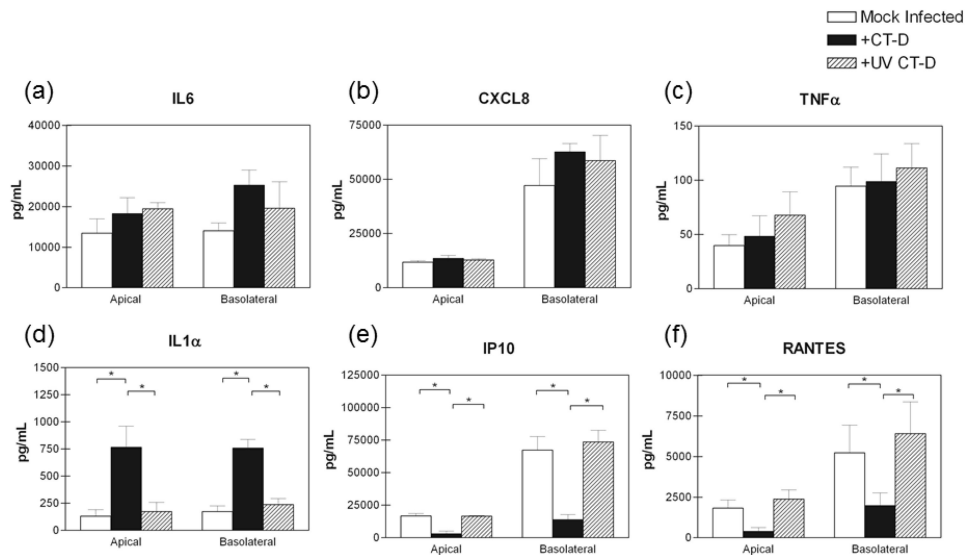


Figure 8. UV inactivated chlamydiae do not elicit a pro-inflammatory cytokine response, but also do not decrease IP10 and RANTES secretion from polA2EN cells

PolA2EN cells were infected with *C. trachomatis* serovar D (>95% infection rate) or exposed to an equivalent amount of UV inactivated chlamydiae. Mock infected polA2EN cells were utilized to ascertain baseline cytokine levels. Apical and basolateral supernatants were collected at 72 hours post-infection or exposure and subjected to cytometric bead array analyses for (a) IL6, (b) CXCL8, (c) TNFα, (d) IL1α, (e) IP10, and (f) RANTES. Cytokine values represent mean values ± SD from 3 independent experiments each performed in triplicate. Statistical analyses between parameters were performed using a One Way ANOVA (** p<0.001).

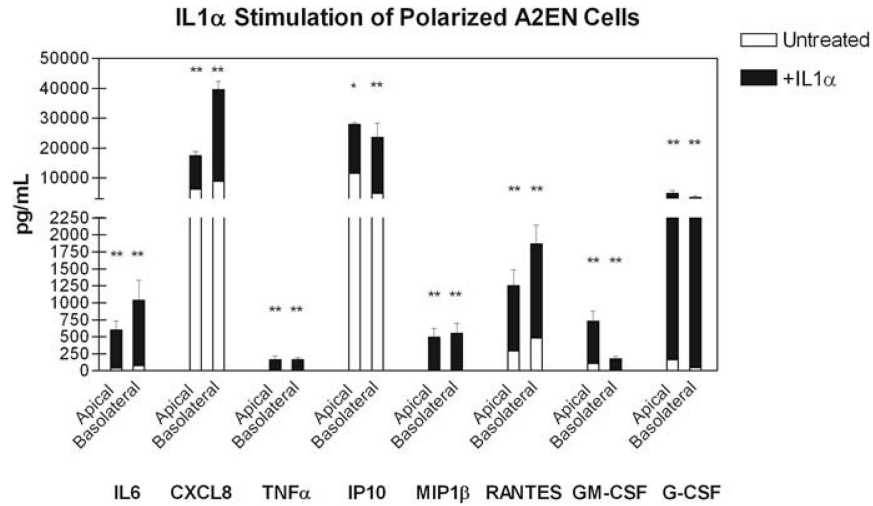


Figure 9. PolA2EN cells respond to IL1α stimulation by up-regulating pro-inflammatory cytokines

PolA2EN cells were exposed to 20ng/mL IL1α for 24 hours. Apical and basolateral supernatants were collected from untreated and treated cells and were subjected to cytometric bead array analyses for IL6, CXCL8, TNFα, IL1α, IP10, and RANTES, GM-CSF, and G-CSF. Cytokine values represent mean values ± SD from 3 independent experiments each performed in triplicate. Statistical analyses between untreated and treated parameters were performed using a One Way ANOVA (* p<0.05, ** p<0.001).

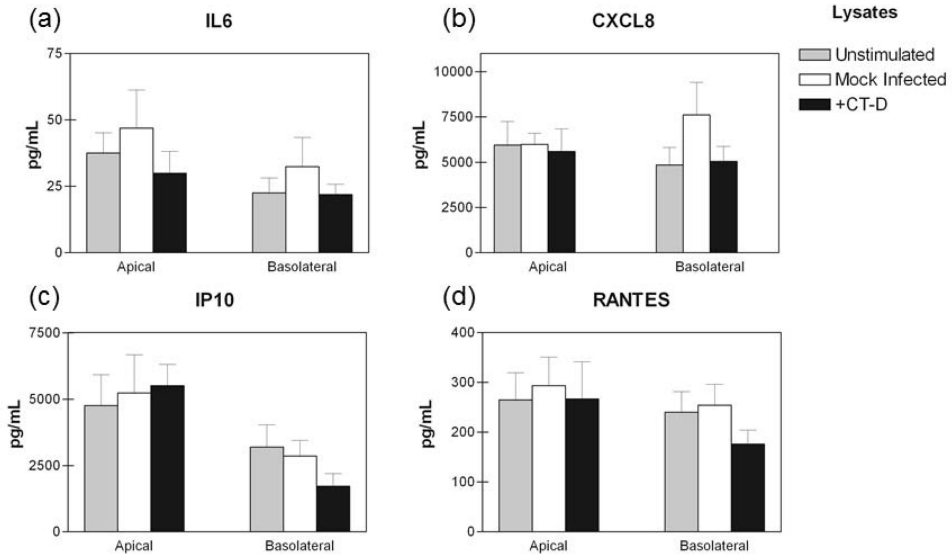


Figure 10. Lysates from *C. trachomatis* infected polA2EN cells do not amplify the pro-inflammatory cytokine response in non-infected cells
 PolA2EN cells were exposed to filtered lysates from mock infected and *C. trachomatis* serovar D infected polA2EN cells for 4 hours. Lysates were removed, and fresh medium was incubated with cells for 12 hours. Unstimulated polA2EN cells were utilized as a control. Apical and basolateral supernatants were collected and subjected to cytometric bead array analyses for (a) IL6, (b) CXCL8, (c) IP10, and (d) RANTES. Cytokine values represent mean values \pm SD from 3 independent experiments each performed in triplicate. Statistical analyses between parameters were performed using a One Way ANOVA.

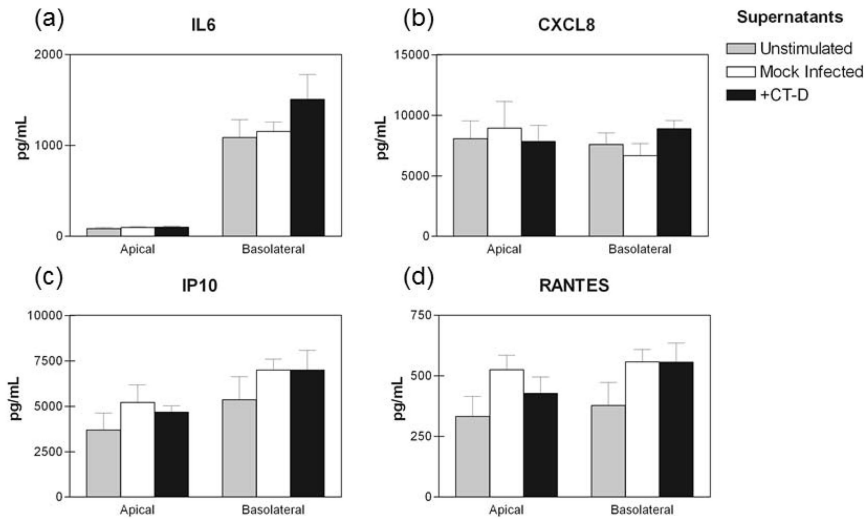


Figure 11. Supernatants from *C. trachomatis* infected polA2EN cells do not amplify the pro-inflammatory cytokine response in non-infected cells

PolA2EN cells were exposed to filtered supernatants from mock infected and *C. trachomatis* serovar D infected polA2EN cells for 4 hours. Supernatants were removed, and fresh medium was incubated with cells for 12 hours. Unstimulated polA2EN cells were utilized as a control. Apical and basolateral supernatants were collected and subjected to cytometric bead array analyses for (a) IL6, (b) CXCL8, (c) IP10, and (d) RANTES. Cytokine values represent mean values \pm SD from 3 independent experiments each performed in triplicate. Statistical analyses between parameters were performed using a One Way ANOVA.